



Journal of Enzyme Inhibition and Medicinal Chemistry

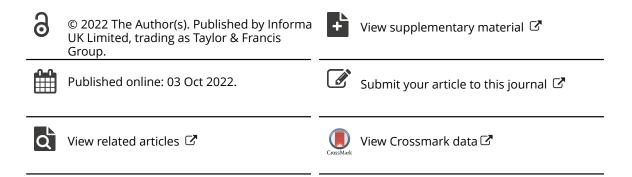
ISSN: (Print) (Online) Journal homepage: https://www.tandfonline.com/loi/ienz20

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To cite this article: Juliusz Maksymilian Walczak, Dorota Iwaszkiewicz-Grześ, Michalina Ziomkowska, Magdalena Śliwka-Kaszyńska, Mateusz Daśko, Piotr Trzonkowski & Grzegorz Cholewiński (2022) Novel amides of mycophenolic acid and some heterocyclic derivatives as immunosuppressive agents, Journal of Enzyme Inhibition and Medicinal Chemistry, 37:1, 2725-2741, DOI: <u>10.1080/14756366.2022.2127701</u>

To link to this article: <u>https://doi.org/10.1080/14756366.2022.2127701</u>



RESEARCH PAPER

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Novel amides of mycophenolic acid and some heterocyclic derivatives as immunosuppressive agents

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ABSTRACT

The group of 18 new amide derivatives of mycophenolic acid (MPA) and selected heterocyclic amines was synthesised as potential immunosuppressive agents functioning as inosine-5'-monophosphate dehydrogenase (IMPDH) uncompetitive inhibitors. The synthesis of 14 of them employed uronium-type activating system (TBTU/HOBt/DIPEA) while 4 of them concerned phosphonic acid anhydride method (T3P/Py) facilitating amides to be obtained in moderate to excellent yields without the need of phenolic group protection. Most of optimised protocols did not require complicated reaction work-ups, including chromatographic, solventconsuming methods. The biological activity assay was performed on the T-Jurkat cell line and peripheral mononuclear blood cells (PBMCs) which are both dedicated for antiproliferative activity determination. Each of designed derivatives was characterised by reduced cytotoxicity and benzoxazole analogue (A2) revealed the most promising activity. Subsequently, an observed structure-activity relationship was discussed.

ARTICLE HISTORY

Received 21 July 2022 Revised 3 September 2022 Accepted 19 September 2022

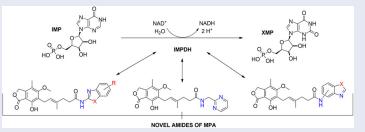
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KEYWORDS

Mycophenolic acid; amide derivatives; heterocycles; benzoxazole; IMPDH inhibition

GRAPHICAL ABSTRACT



MOST WIEDZY

Introduction

Inosine-5'-monophosphate dehydrogenase (IMPDH) is an enzyme responsible for oxidation of inosine-5'-monophosphate into xanthine-5'-monophosphate (XMP) exploiting water molecule as well as NAD⁺ redox potential. This biotransformation is crucial for proper cell growth due to the fact that it triggers guanyl nucleotides formation which serve as RNA and DNA precursors, the energy reservoir for translation process, the cofactor for G-proteins as well as glycosylation precursor. Depletion of guanine-based nucleotides, which is induced by redirecting biosynthesis into adenine-based nucleotides pathway, results in impairment of de novo and salvage route of purine and pyrimidine nucleotides biosynthesis. This misregulation proceeds from phosphoribosyl pyrophosphate synthetase (PRPP) and ribonucleotides reductase (RNR) stimulation dependence accruing from guanine and adenine nucleotides. When the previous one activates enzymes' bioactivities, the latter one inhibits them. Eventually, enhancement of adenosine nucleotides pool effects in malfunctioning of rapidly proliferating human cells¹.

Mycophenolic acid (MPA) is one of the uncompetitive and reversible inhibitors of IMPDH. It means that MPA and IMP bind to an enzyme simultaneously thus blocking further biosynthesis of guanine-based nucleotides causing significant effects on the cell's biochemistry¹⁻³. Thereupon, MPA was previously found to inhibit human lymphocytes G and T responses to mitogenic stimulation thus resulting in immunosuppressive activity observation. This feature led to new immunosuppressants establishment, sodium mycophenolate (MMS) and mycophenolate mofetil (MMF) namely⁴. Due to their antiproliferative properties, they have been put to use in acute graft rejection prophylaxis and psoriasis. Despite severe side-effects (e.g. secondary microbial infections, red blood cells activity) arising from MPA usage, it is still in use for its unique risk-reducing feature of graft loss and acute rejection outperforming other immunosuppressants, e.g. Azathioprine^{1,5,6}. It is worth mentioning that some attempts towards MPA-based

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B Supplemental data for this article is available online at https://doi.org/10.1080/14756366.2022.2127701

bioactive compounds isolation from natural resources are still being made thus showing not only synthetically approachable derivatives. Some of them originate from marine-derived fungus *Penicillium sp.* SCSIO sof101 and *Rhizopsus oryzae* which are a reservoir for penicacids that exhibit from moderate to weak antibacterial properties^{7,8}. Targeting of IMPDH by the MPA molecule may lead to anticancer properties exhibition *via* other molecular mechanisms. Enzyme's inhibition can lead to a decrease in the expression of several antiapoptotic proteins and the deactivation of caspases which is an excellent starting point in osteosarcoma chemotherapy and the antimetastatic drugs designing in general⁹, anticancer potential of MPA is still under investigation^{10,11}.

Bioactivity of MPA is preserved due to the selected functionalities present within its structure. Lactone and aromatic moiety, free phenolic group,/E/-shaped double bond, and carboxylic acid moiety are required to maintain compound's activity thus making radical structural modifications forbidden^{1,4,12}. Nevertheless, some amidetype derivatives were synthesised and positively biologically evaluated. Anilide-type derivatives of MPA have shown good bioactivity towards human and protozoal IMPDH making them potential immunosuppressants and antiprotozoal agents¹³. Hydroxamic acid and aminated heterocyclic derivers of MPA are prone to inhibit IMPDH as well as histone deacetylase (HDAC) rendering them for being anticancer agents for influencing some of the expression level of anticancer genes in abnormal cells, beside of its antiproliferative properties^{14,15}. Another example of a unique amide-like derivative of MPA is connected with MPA-quinic acid conjugate. It bears subtly smaller activity towards IMPDH and protective effect against apoptosis at once. It also improved insulin release profile in INS-1E cell lines thus showing how such a bioactive conjugate would shift cytotoxicity at all¹⁶. Mycophenolic adenine dinucleotide (MAD) analogues were rationalised as another inclined IMPDH inhibitors¹⁷. MPA and biogenic amino acids showed moderate antibacterial properties, especially towards S. aureus and K. pneumoniae, most likely due to the inhibition of bacterial IMPDH¹⁸. Benzylderived amides of MPA are thought to be potent anticancer drugs candidates due to similar activity towards IMPDH as MPA and unique inhibitory properties in relation to breast, prostate, and glioblastoma cell lines¹⁹. Additionally, some amide derivers of MPA, especially one bearing long aliphatic spacer and terminal imidazole structural motif, tend to impede Toxoplasma gondii growth better than referential MPA molecule, both in in-vitro and in-vivo conditions, thus presenting excellent anti-protozoal properties and making it a potential drug candidate²⁰.

Noteworthy, IMPDH inhibition concerns wide class of compounds, e.g. structures possessing urea moieties or various heterocyclic units, which have been investigated towards immunosuppressive, anticancer, antibacterial, and antiviral activities^{1,21–23}.

Heterocyclic motifs are one of the most important constitutional patterns among all existing pharmaceuticals' structures. In 2014, four of five most common prescribed drugs in the US contained heterocyclic fragments²⁴. Of 59% of small-molecule remedies sold in the US, in the same year, also carried heterocyclic designs, but this time it belonged to the nitrogen-based heterocyclic compounds family²⁵. These facts render heterocycles as being the most privileged and significant molecular architectures in the drug designing area. Heterocyclic structural units are known for their remarkable physicochemical properties. They may serve as versatile bioisosteres as they can be tuned in their lipophilicity, polarity and water-solubility upon currently desirable properties in molecular targets reaching. These features may be manipulated in relation to their aromaticity and polar properties developed by the presence of strong electronegative atoms²⁴. Referring to previously mentioned examples, heterocyclic structural motifs such as 1,2,4-triazole, incorporated into MPA molecule, have shown similar activity towards IMPDH as MPA within itself¹⁵. Inhibitory properties of benzoxazole-based derivatives towards parasitical IMPDH have also been proved, making them potential antiparasitic agents²⁶. Heterocyclic compounds are also known for their anti-inflammatory²⁷, anticancer^{28,29}, antimalarial³⁰, and fungicidal properties³¹ and these parameters may be a source of added value beside antiproliferative nature of MPA.

In this work, we submitted molecular docking assay depicting binding force in substrate-enzyme complex between novel amide derivatives of MPA and IMPDH, then the synthesis of novel MPA derivatives and their structural characterisation and eventually biological evaluation involving T-Jurkat cell lines as well as peripheral mononuclear blood cells (PMBCs). MPA core was chosen upon previous literature premises¹⁻⁴ and the significant role it plays in graft rejection prophylaxis, and more precisely, the significant cytotoxicity it shows to healthy cells⁵. Aforementioned examples of biological utility of amide-like derivatives of MPA¹³⁻²⁰ prompted us to explore another class of amide derivers with heterocyclic structural motifs incorporated. The latter pattern utilisation was justified by its unique physicochemical properties, common application in small-molecule drugs^{24,25} and plenty of bioactivities^{15,26–33} which may serve as an added value to the referential MPA molecule. All of these elements brought together may represent attractive novel antiproliferative compounds in medicinal chemistry by lowering cytotoxic effects on healthy cells and maintaining approximate antiproliferative properties in relation to MPA-based drugs (MMS, MMF, or MPA within itself) at the same time.

Results and discussion

Molecular docking

To verify that compounds **A1–A18** are able to effectively bind to the IMPDH enzyme's active site, molecular docking studies were performed. The X-ray structure of IMPDH enzyme was retrieved from the Protein Data Bank (Protein Data Bank accession code 1JR1) and properly prepared for docking calculations. The docking procedure of the optimised ligands was performed using AutoDock Vina version 1.1.2 software (Molecular Graphics Laboratory, The Scripps Research Institute, LaJolla, CA). The calculated results for the proposed structures of compounds A1–A18 indicated that they could, at least theoretically, efficiently bind to the active site of IMPDH protein. The free energies of binding of examined compounds were at the satisfactory

Table 1. Fre	e energies o	f binding	calculated fo	r compounds	A1–A18 and MPA.
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Entry	Free energy of binding [kcal·mol ⁻¹]		
MPA	-7.8		
A1	-9.3		
A2	-9.1		
A3	-9.4		
A4	-9.5		
A5	-8.9		
A6	-9.5		
A7	-8.7		
A8	-8.8		
A9	-9.2		
A10	-9.4		
A11	-9.1		
A12	-9.1		
A13	-9.0		
A14	-9.1		
A15	-9.2		
A16	-9.0		
A17	-9.2		
A18	-9.1		

level in the narrow range of -8.7 to -9.5 kcal·mol⁻¹ (Table 1) and were noticeably lower than the free energy of binding of MPA (-7.8 kcal·mol⁻¹) used as reference. The most favourable free energies of binding were determined for compounds A4 and A6 (-9.5 kcal·mol⁻¹ in both cases).

Aforementioned results show that the influence of heteroaromatic core, loci of amino group as well as other functionalities on the free energy of binding with IMPDH can be distinguished. Within various heteroaromatic compounds, benzimidazole-based derivative (A3) turned out to bound most strongly to an enzyme, followed by benzothiazole (A1), benzoxazole (A2) and finally - pyrimidine counterpart (A7). Within the amino group substitution variety, one can see the strongest enzyme-affinity of 6-aminated benzothiazole (A4), then 2and 5-aminated equivalents (respectively: A1 and A5). In the case of benzoxazole motif, 5-aminated derivative (A6) came out as more enzyme-binding effective than 2-aminated analogue (A2). Considering the structure-activity relationship for a class of novel benzothiazole derivatives (A8-A18), one can observe no clear-cut differences in the value of binding energy towards IMPDH. However, dimethylated (A10) and nitro (A15) derivatives have the highest affinity for the enzyme. Moreover, this comparison showed that the halogen derivers (A11-A13 and A18) do not differ much from each other, even taking trifluoromethylated amide (A14) into this consideration. Additionally, 4substituted compounds (A16-A18) did not differ from their 6-substituted counterparts (A8, A9, and A12) in case of the enzyme affinity except for the methoxy derivatives (A8 and A16).

The visualisation of the putative binding mode for the representative compound A2 $(-9.1 \text{ kcal} \cdot \text{mol}^{-1})$, which demonstrated the most promising activity in the biological evaluation, is shown in Figure 1. It can be noticed that compound A2 is very well accommodated in the IMPDH active site and is surrounded by several important amino acid residues. For example, the mycophenolic part of the compound is in the close distance to the Gly326 (3.42 and 3.54 Å), Thr333 (2.70 Å), and Gln441 (3.22 Å) amino acid residues suggesting the presence of electrostatic interactions. Furthermore, the π - π stacking interaction (3.66 Å) with IMP molecules was detected. In addition, the benzoxazolyl residue of compound A2 may also electrostatically interact with several amino acid residues, e.g. Ser276 (2.71, 2.93, and 3.27 Å), and Gln441 (2.99 and 3.62 Å). All of these plausible interactions may stabilise the enzyme-ligand complex ensuring appropriate accommodation of compound A2 and higher inhibitory activity against target protein.

It can be seen that structure-activity relationship for the new amide derivatives of MPA is worth establishing upon biological testing, mostly due to differing free energy binding values and to see if heteroatom affects specific interactions with enzyme subunits thus facilitating further elaborations of novel immunosuppressants class, especially those based on benzothiazole motif, which turned out to be the most accessible for us and promising upon *in-silico* studies.

Chemistry

The newly devised compounds A1-A18 were obtained via typical reaction between carboxylic acid and amine moiety in the electrophilic activation of carboxylic acid functional group manner. Doubly activated MPA in the TBTU/HOBt system, in the presence of tertiary amine, was suggested to be more amine-accessible than in the standard system since additive presence (HOBt) enhances reactivity and reduces unreactive intermediate formation³⁴. T3P-based method was used in the case of amino acid derivatives¹⁸ and then successfully applied for amines bearing its amine group in non-heteroaromatic region. This selectivity towards activating agent was probably due to the lack of nucleophilicity of heterocyclic amines towards activated carboxylic acid species, potential spontaneous heterocyclic tautomerisation within 2-amino-1,3-azole unit and intermediate state destabilisation resulting from immediate vicinity of heteroatoms. Albeit it needs to be thoroughly investigated since 2-aminomethylpyridines did not give desired products via T3P activation thus indicating other factors importance (e.g. solubility of substrates or pH imbalance). Chloroformate/tertiary amine system worked in Method A conditions however resulting in significantly lower yields, what has not been presented in this work. A general presentation of conducted syntheses and their outcomes is given in Scheme 1 for Method A and Scheme 2 for Method B.

Acetonitrile (ACN) seems to be the most promising solvent (for A1, A2, A8–A18) for its polar and aprotic properties, volatility, and non-adsorptive character. Acetone is prone to form imine species, DMF has a tendency for lingering at products' surface while dimethyl sulfoxide (DMSO) causes removal difficulties. Non-polar solvents such as toluene, dichloromethane, or ethyl acetate (AcOEt) did not work at all, whereas polar solvents usage is at odds with electrophilicly-activated intermediates reaction

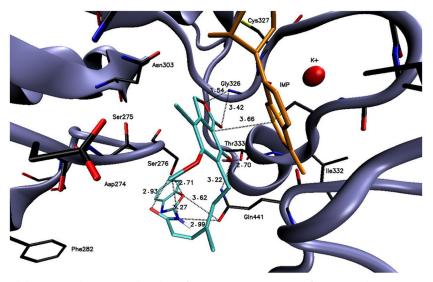
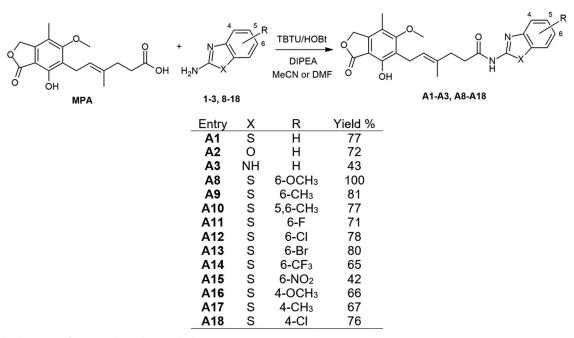
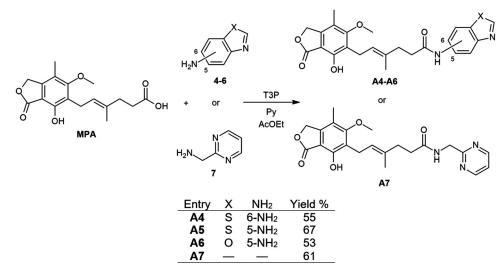


Figure 1. Docked binding mode and distances to the amino acid residues of IMPDH enzyme's active site for compound A2.



Scheme 1. Amide derivatives of MPA synthesised via Method A.



Scheme 2. Amide derivatives of MPA synthesised through Method B.

mechanisms. For **A4–A7** cases, AcOEt worked, probably as it is well-suited for stabilising intermediate species established between non-heteroaromatic amine and **T3P**. Dichloromethane and ACN did not work during reaction's optimisation and just like before – polar solvents are not applicable.

Most of compounds under investigation were isolated without chromatography methods. Resulting derivatives of **MPA**, namely **A1**, **A2**, **A8**–**A18**, were generally poorly soluble in ACN and hence were daringly washed with cold solvents to get rid of impurities, unreacted activating agents, and tertiary amine as well. However, some dissimilarities were observed in case of 4-substitued derivatives (**A16–A18**) and 2-aminobenzoxazole deriver (**A2**) regarding to 6-substituted ones. That tendency occurred probably due to different influence of substituent/heteroatom towards hydrogen bond formation thus increasing solubility in both polar and apolar solvents. Another exception was observed in **A3** case as DMF was used for enhancing originate amine solubility. This solvent was also used for the washing step which resulted in difficult to remove, solvent residual peak presence in the ¹H NMR (nuclear magnetic resonance spectroscopy) spectrum. Examples **A4–A7** possess completely different solubility trends thereout different methods and solvents were involved for preparing them.

For A1, A8, A11, and A14 cases, 2.0 equivalents of tertiary amine (DIPEA) were applied due to standard procedure present in the literature³⁵. However, in most cases only 1.0 equivalent was utilised on account of reaction mechanism. Initially, each reaction was carried out in a manner consistent with the reaction mechanism, but the one proposed in the literature was also studied. Differences in the base amounts may happen due to the high stability of salt formed upon MPA and amine dissolution. Increasing the amount of DIPEA alters pH balance which affects the tendency of amine to react with activated MPA as it may less readily occur in protonated form. Base excess is believed to increase coupling rates and to disrupt hydrogen bonds responsible for conformation stabilisation which affects the availability of amino groups³⁶. A3 was synthesised with individual quantities of base and activating

Correlation between the character of substituent in 2-aminobenzo[d]-1,3-diazole and its condensation yield with MPA is as follows: electron withdrawing group has a potential upon diminishing reaction's yield as it is clearly observed for A14 and A15 and electron donating group improves reaction's outcomes regarding A8-A10. 6halogenated derivatives, namely: A11, A12, and A13, should present quantitively worsened results as inductive withdrawing effect ought to lower nucleophilicity of amino group, however yields in this case were rather noncongruent with this tendency and showed decreased reaction outcomes with halogen electronegativity rise, A13, A12, and A11 sequentially. Electronegativity impact was also observed in A1-A3 cases since its growth developed lowered yields, nevertheless heterocyclic tautomerisation within amine's structure cannot be neglected. Unclear correlation was noticed in 4-substituted cases, namely: A16, A17, and A18, as they present opposite trend. It probably results from different impact of substituent position on hydrogen bonding thus making solubility the limiting factor of reactions outcomes. Pretty convergent yields can be seen for A4-A7 cases. They represent the class of non-heterocyclic amines which are characterised by different reactivity tendencies than previously mentioned examples. Aliphatic amine (A7) has been most successfully conjugated with MPA in this subgroup of derivatives as indirect impact of aromatic system does not decline nucleophilicity of amino group. In all cases, the solubility of reactants may influence reaction's outcomes acting as both the driving force and limiting factor of reaction. Considerations on the effect of substituent nature on melting point³⁷ are presented in Electronic Supplementary Information (ESI).

MPA as a fully substituted aromatic compound possesses no protons in the aromatic region. This fact simplifies amides' ¹H NMR spectrum analyses since newly incorporated moieties contribute aromatic and amide protons solely. All spectra are determined by the deshielded amide proton and phenolic one as well as recently introduced aromatic species. Remaining signals represent MPA structure and are sometimes shifted due to magnetic environment modifications. Some lapping may occur as implemented amines bear functional groups or spacers appearing in the same ¹H NMR regions as MPA structural units. ¹³C NMR spectra are also characterised by MPA core and added value in the shape of newly introduced carbon atoms. Some dissimilarities in the nucleus characteristics may occur (especially in the case of amide group carbon) as well as for amine-based units. The latter one is represented by deshielded C(2) heteroaromatic carbon and C(3a)/C(7a) positions. Remaining loci differ with heteroatom electronegativity and substituent character, sometimes precluding precise carbon atoms assignment. Some ¹H and ¹³C NMR spectra show long and short distance spin decoupling brought by fluorine atoms, namely in A11 and A14 cases. More detailed information about the character of heteroaromatic interactions within ¹³C NMR spectroscopy may be found in the literature^{38,39}.

Present coupling reaction between MPA molecule and selected heteroaromatic amines provides a convenient method of amide bond synthesis since MPA is a multifunctional, synthetically challenging compound. Beforehand, typical synthesis exploiting the MPA molecule demanded phenolic group protection^{15,40} as it may

hamper nucleophilic potential by impairing possible interference with activated carboxylic acid moiety of other molecules. Our work delivers simple, undemanding and solvently balanced synthetic protocol which in most cases does not require chromatography purification methods and more importantly, does not involve phenolic group protection/deprotection manner being in line with current mode to the synthesis of MPA derivers^{41,42}. Moderate to excellent yields and spectrally traceable products' make this approach remarkably potent and applicable in the synthesis of heterocyclic-derived amides of MPA.

Biological investigations

Drug toxicity that affects the cells can cause a variety of serious side effects, including uncontrolled suppression of the immune system. In order to check the cytotoxicity of the tested compounds, the XTT test was performed, the results of which are presented in Table 2. All tested compounds were characterised by lower cytotoxicity than native MPA. Unfortunately, several compounds (marked with \downarrow) were partially insoluble in culture despite surfactant addition. They were investigated too, but one could not use them as an immunosuppressive drug in their current form. Taking into account the EC₅₀ values obtained in the proliferation tests (VPD540 staining, Table 3), the compound **A2** was selected as the one with the highest immunosuppressive potential and the possibility of its later use in therapy, and is presented in the graphs (Figures 2–4).

In order to test the immunosuppressive effect of MPA derivatives, we decided to use T-Jurkat cell line as a model of human lymphocytes and PBMCs which are widely used in research and toxicology applications as *in vitro* model of the immune response, and provides answers to questions about the toxicity of potential new drug compounds.

Determination of cytotoxic activity using XTT method

Table 2 is collected IC_{50} values of tested derivatives on PBMCs and T-Jurkat cells under XTT assay conditions. In the case of the PBMC approach, each derivative exhibits lower cytotoxic properties than reference (MPA). Similar conclusions may be drawn from assay performed on T-Jurkat cell line albeit **A2** possesses approximate IC_{50} value to MPA. Due to the limited solubility of **A8**, **A9**, **A11**, **A12**, **A15**, **A16**, and **A17** one could not receive reliable results for PBMCs. In the T-Jurkat-based method, some IC_{50} values were not obtained and thus cell viability was measured at the

Table 2. IC_{50} [µM] values of derivatives of MPA based on XTT test.

	PBMCs		T-Jurkat	
Entry	IC ₅₀ [μM]	Viability [%]	IC ₅₀ [μM]	Viability [%]
MPA	10.522 ± 0.015	-	0.509 ± 0.008	_
A1	$74.570 \pm 0.040^{*}$	_	-	68
A2	15.923 ± 0.009	_	1.326 ± 0.002	_
A3	94.538 ± 0.047	_	-	74
A4	24.639 ± 0.015	-	37.789 ± 0.002	-
A5	45.730 ± 0.023	-	75.040 ± 0.005	-
A6	55.487 ± 0.028	-	31.407 ± 0.002	-
A7	70.480 ± 0.035	-	10.739 ± 0.003	-
A8	93.196±0.047 (↓)	-	51.124 ± 0.008	-
A9	94.600±0.047 (↓)	-	85.700 ± 0.004	-
A10	129.408 ± 0.064	-	-	51
A11	139.858 \pm 0.070 * (\downarrow)	-	-	56
A12	62.975±0.031* (↓)	-	$12.833 \pm 0.008^{*}$	-
A13	$63.173 \pm 0.032^{*}$	-	58.069 ± 0.002	-
A14	$81.482 \pm 0.041^*$	-	34.749 ± 0.003	-
A15	137.633±0.069* (↓)	-	12.717 ± 0.005	-
A16	54.578±0.027 (↓)	-	-	69
A17	- (↓)	65	-	82
A18	$74.426 \pm 0.037^{*}$	-	10.261 ± 0.001	-

Significance was calculated using the *T*-test; significant results are marked with *(p < 0.05) with MPA as a control; (1) – precipitating compound.

Table 3. EC_{50} $[\mu M]$ values of derivatives of MPA based on dye-based proliferation assay VPD450.

	PBMCs		T-Jurkat	
Entry	EC ₅₀ [μM]	Viability [%]	EC ₅₀ [μM]	Viability [%]
MPA	0.723 ± 0.011	-	0.490 ± 0.002	-
A1	-	82	_	_
A2	5.601 ± 0.022	-	0.850 ± 0.004	_
A3	119.600 ± NAN**	-	-	55
A4	62.260 ± 0.010	-	14.730 ± 0.074	-
A5	$76.980 \pm 0.010^{*}$	-	36.050 ± 0.180	-
A6	58.270 ± 0.021	-	13.840 ± 0.069	-
A7	29.450 ± 0.025	-	9.510 ± 0.047	-
A8	133.500 ± 0.007*(↓)	-	22.040 ± 0.110	-
A9	136.200 ± 0.011(↓)	-	22.720 ± 0.113	-
A10	$85.240 \pm 0.012^{**}$	-	26.690 ± 0.133	-
A11	83.070±0.012**(↓)	-	63.210 ± 0.316	-
A12	38.370 ± 0.017(↓)	-	12.160 ± 0.061	-
A13	39.470 ± 0.013	-	12.640 ± 0.063	-
A14	30.710 ± 0.008	-	12.580 ± 0.063	-
A15	44.590 ± 0.013(↓)	-	13.370 ± 0.067	_
A16	46.290 ± 0.011(↓)	-	-	73
A17	82.140 ± NAN***(↓)	-	-	69
A18	14.700 ± 0.020	-	11.280 ± 0.056	-

Significance was calculated using the T-test; significant results are marked with *(p < 0.05), **(p < 0.01), or ***(p < 0.001) with MPA as a control; NAN – not a number; (\downarrow) – precipitating compound.

highest concentration using trypan blue exclusion (for A1, A3, A10, A11, A16, and A17 namely). Even despite the impossibility of accurate results showing in each case, the general tendency shows that the incorporation of benzothiazole, benzoxazole, benzimidazole, and pyrimidine systems decrease cytotoxicity and in terms of precise values it is usually several dozen times lowered comparing to the reference compound. The cytotoxicity was influenced by amino group position in heterocyclic motifs. Entries A1, A4, and A5 show that 6-aminated benzothiazole affects cytotoxicity stronger than 5- and 2-aminated ones in succession. Similar effects may be seen in entries A2 and A6 for aminated benzoxazole. Correlation between the character of substituent and cytotoxic effect is unclear, however these possessing electron-rich substituents seem to be more effective than the neutral ones (A13, A14, and A18 versus A10). The tendency of IC₅₀ values in T-Jurkat cells can be determined similarly, however 6-methoxy (A8) as well as 6-nitro (A15) and 6-methyl (A9) derivatives show: the more electron-rich group is attached, the stronger the cytotoxic effect of the corresponding derivative. Halogen impact may be noticed in this case as in the group A12-A14 the increase of electronegativity builds cytotoxicity up. Little difference between 6substituted and 4-substituted derivatives (A12 and A18) may be noted.

Eventually, **A2** is the most invasive derivative as it affects both PBMCs and T-Jurkat cells, however still weaker than the reference compound. Remaining amides are clearly less toxic than MPA and some correlations between the type of substituent, loci of amino group and cytotoxicity may be observed. Electron-rich moieties seem to enhance derivers' toxicity towards exploited cells, albeit this relation is unclear since limited by solubility of tested amides in conditions given.

Figure 2 presents the response curve (concentration of tested compound is assigned to viability of cells) prepared for a few different concentrations of MPA and **A2** in a suitable medium. On the basis of these measurements one can determine optimal IC_{50} value for referential MPA and the most promising compound **A2**. As it was mentioned before, **A2** possesses lowered cytotoxicity towards human and T-Jurkat cells which is represented by this graph. In general, plots are quite convergent in the lowest and

highest concentrations and differ evidently in the medium range. Noteworthy, amide **A2** indicated the highest antiproliferative activity within the designed compounds (Table 3) together with lower cytotoxicity in comparison to parent MPA (Table 2).

Effect of MPA and A2 on apoptosis

The apoptotic effects of MPA and A2 were evaluated by flow cytometry after staining with Annexin V and Pl. Annexin V can be detected in early and late stages of apoptosis, whereas PI can only be detected in late apoptosis or necrosis. Due to the level of necrotic cells at about 1%, it was not taken into account during the analysis. As shown in Figure 3, treatment with compound A2 and MPA as a control for 96 h, late apoptosis cells could be observed mainly when the cells were treated with a concentration close to the EC₅₀ value (0.723 μ M for MPA and 5.601 μ M for A2). In the case of MPA in a concentration of $10\,\mu$ M, one can observe the presence of over 13% of late apoptosis cells, but only about 4% of early apoptosis cells. This situation changes significantly in the case of a 10-fold lower concentration, where over 17% of early apoptosis cells and over 11% of late apoptosis can be found. Derivative A2 showed a similar effect, however, at a concentration of 10 µM, where one can see more than 14% of early and late apoptosis cells. It also indicated that the inhibition of cell proliferation, even at the $10 \,\mu$ M concentrations (Figure 3(B)), was mainly due to the influence of tested compounds.

Determination of antiproliferative activity using Violet Proliferation Dye (VPD) 450 in flow cytometry

The antiproliferative effects of MPA and tested compounds were evaluated by flow cytometry after staining with VPD450 which is one of the recommended dyes, characterised by a relatively low toxicity and the possibility of observing cell division through the loss of a fluorescent dye⁴³. The representative dose-response curve to determine the EC_{50} of MPA and A2 are presented at Figure 4 and exemplary histograms with appropriate microscopic pictures are provided in ESI (ESI supplementary Figure 1). As it can be observed in Table 3, all compounds have an EC₅₀ value greater than MPA for both PBMCs and T-Jurkat cells. Compound with the most activity is A2 with an EC_{50} value 5.601 μ M. However, taking into account that this compound is characterised by lower cytotoxicity than MPA, it may constitute a suitable candidate for future research. Additionally, there was no change in cell proliferation when they were treated with DMSO and the solubilisation additive, suggesting that the observed reduction of proliferation was exclusively due to the presence of MPA or tested compounds in the cultures.

Results presented in Table 3 show similar tendencies as in Table 2. Unambiguous outcomes may be observed by comparing PBMCs and T-Jurkat cells' behaviour towards tested compounds as they exhibit convergent trends. For the first conditions, one can notice loci differentiation in case of 6- and 5-aminated benzothiazole-derived amides (namely A4 and A5) for the first instance presenting higher antiproliferative properties than the second one. Under these circumstances, 2-aminated deriver's properties (A1) could have not been determined under PBMC culture medium. A5 shows worsened effectivity in comparison to A2 (that strongly affects human cells) thus exposing again amino group loci importance at bioactivity matter. In terms of the electronic nature of substituents, one can discern a clear-cut difference between 6halogenated derivatives of benzothiazole and remaining ones. 6chloro (A12), 6-bromo (A13), and 6-trifluoromethyl (A14) cases

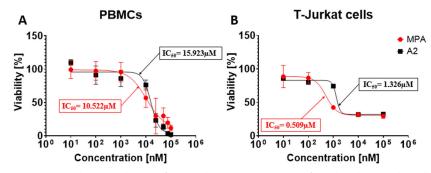


Figure 2. Representative dose-response curve to determine the IC_{50} of MPA and **A2**. (A) Human PBMCs for 96 h and (B) T-Jurkat cell line for 48 h were cultured in the presence of different concentrations (100; 75; 50; 25; 10; 1; 0.1 and 0.01 μ M) of MPA (red) and **A2** (black). Next, plates were incubated for 24 h with XTT reagent. The conversion of water-soluble yellow tetrazolium XTT salt into orange formazan was monitored by measuring the optical density at 450 nm on microplate spectrophotometer. For XTT assay at least three experiments were performed and values are presented as mean ± SD.

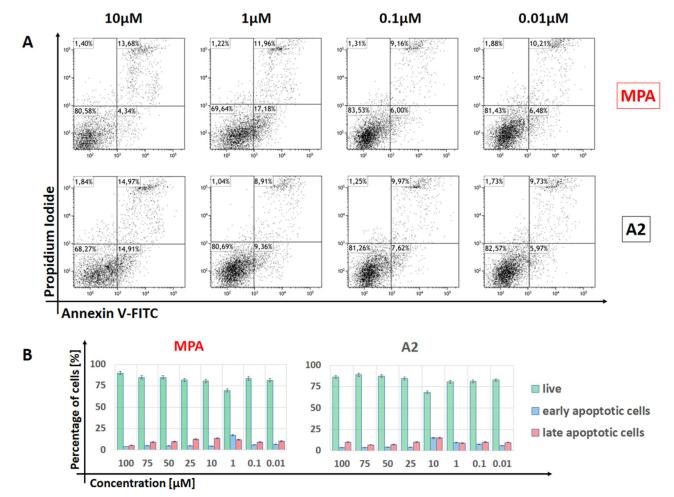


Figure 3. Detection of cell apoptosis in PBMCs by Annexin V-PI staining assay using flow cytometry. (A) Representative scatter plots of cells treated with different concentrations (10; 1; 0.1 and 0.01 μ M) of MPA and **A2** for 72 h and stained with Annexin V-FITC and propidium iodide (PI). (B) Cells divided into three groups according to Annexin V/PI results: living cells (Annexin – PI–); early apoptotic cells (Annexin + PI–) and late apoptotic cells (Annexin + PI+).

manifest significantly increased inhibitory properties towards PBMCs than 6-methoxy (**A8**) and 6-methyl (**A9**) examples, whereas 6-fluoro (**A11**) does not fit into this trend revealing poor antiproliferative features. Loci of chlorine atom or methoxy group may also affect bioactivity as 4-chloro (**A18**) and 4-methoxy (**A16**) derivatives present good affinity towards cell division rupture (see 6-chlorinated and 6-methoxylated counterparts, **A12** and **A8**, respectively). T-Jurkat cell line assay reveals the same leaning,

nevertheless displaying lowered EC_{50} values for 6-methoxy (**A8**), 6-methyl (**A9**), 5,6-dimethyl (**A10**), and 6-nitro (**A15**) derivers.

Ultimately, these biological studies have shown that none of the synthesised amides exhibit antiproliferative properties greater than reference compound, however one of them – **A2**, reveals close enough value of EC_{50} to MPA in both tests. Further 2-aminobenzoxazole-derived compounds' functionalisation need to be proceeded and bioactivity of resulting derivers should be

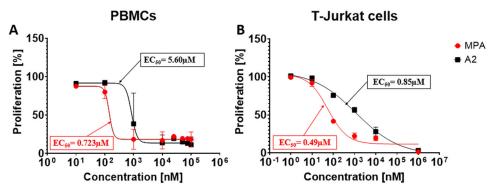


Figure 4. Representative dose-response curve to determine the EC_{50} of MPA and A2. (A) VPD 450-labelled human PBMCs were cultured in the presence of different concentrations (100; 75; 50; 25; 10; 1; 0.1 and 0.01 μ M) of MPA (red) and A2 (black) and stimulated with magnetic beads coated with anti-CD3 and anti-CD28 antibodies for 72 h. (B) VPD 450-labelled T-Jurkat cells were cultured in the presence of different concentrations (100; 10; 1; 0.1 and 0.01 μ M) of MPA (red) and A2 (black) for 48 h. Cell proliferation was analysed using flow cytometry. The results are expressed as the mean ± SD.

performed. Structure–activity relationship has shown unclear correlation, though more electron-rich substituents tend to affect EC_{50} value stronger than neutral ones, there again fluorinated amide does not incorporate into this trend. Rapid change in EC_{50} values ratio for **A8:A12** pair for PBMCs and for T-Jurkat cells approach, from 3.48 to 1.81, reveals obvious differences in tests' complexity as compounds may be shielded by medium components.

Comparison of antiproliferative activity measurements of MPA and **A2** was presented in Figure 4. Plots convergence is slightly inferior than this formed for IC_{50} value figure, especially in the smallest and highest concentrations range. In case of T-Jurkat cells, half maximal effective concentration is more concurrent than for PBMCs. PBMC proliferation measurements done for **A2** and MPA reflect on highly similar antiproliferative properties as staining intensity of PBMCs and cell division's histograms are almost identical (see ESI supplementary Figure 1). It may have positive repercussions for further 2-aminobenzoxazole and MPA derivatives biological evaluations.

The reported compounds have shown eminently lowered cytotoxic effects on both PBMCs and T-Jurkat cells while at the same time exhibiting weaker antiproliferative properties than referential MPA. Despite limited solubility, measurements were generally possible with surfactant addition. Importantly, Tween 20 does not influence the bioactivity at all. Structure–activity relationship for these compounds is unclear, however electron-rich substituents tend to both increase cytotoxicity and antiproliferative properties as well. Fluorine atom does not fit into this correlation, probably due to specific substrate-enzyme interactions occurrence. Loci differentiation may be observed especially in varying amino group position, albeit 4-substituted 2-aminobenzothiazoles seem to affect cell lines stronger than its 6-substituent on bioactivity.

Subsequently, we verified if designed MPA derivatives act as IMPDH inhibitors. According to reported experiment⁴⁴, suppression of cell proliferation caused by IMPDH inhibitors in the presence of guanosine source, e.g. gunosine monophosphate (GMP) is reversible. To this test were selected MPA (reference), **A2** as the most active target amide (Figure 5), and **A18** (active among benzothiazoles), and **A7** as possessing another heterocyclic ring (see ESI supplementary Figure 2). The both Figures show the proliferative activity of the tested compounds against PBMCs at the appropriate concentrations. The mechanism of action of the derivatives was consistent with that of MPA. In the concentration range where inhibition of proliferation occurs without the addition of

GMP (for MPA: 100; 10 and 1 μ M, and for **A2**: 100 and 10 μ M), it can be seen that proliferation increases after the addition of GMP.

In the end, A2 has shown the strongest antiproliferative properties for exhibiting similar properties as MPA. Its lowered cytotoxicity and better solubility can be an added value for determining new 2-aminobenzoxazole and MPA derivatives properties. Its similarity in PBMCs proliferative response to the referential compound renders it into being a potent immunosuppressant, albeit further investigations must be proceeded. The presence of nitrogen and oxygen atoms in newly incorporated amine fragment in A2 may indicate the influence of these elements on the bioactivity of new MPA derivatives since it is present in MMF - the most common and wide-spread drug used in graft rejection prophylaxis. In the sulphur atom case (low cytotoxic 2-aminobenzothiazole derivatives), its presence significantly decreases antiproliferative properties, however, it may affect further biological assays in the context of zinc-based enzymes, similarly to reported in literature HDAC inhibitors^{45–47}.

Summary

The series of 18 novel amide derivatives of MPA were designed for their affinity towards IMPDH in the enzyme's active site, employing in-silico assessment. The synthesis was optimised with uronium (TBTU/HOBt) or phosphonic acid anhydride (T3P) electrophilic activation systems, for carboxylic acid moiety, which are widely available and easy accessible, and were well adjusted for aliphatic, aromatic, and heteroaromatic amines to give target amides in moderate to excellent yields mostly without the need of chromatographic purification. Subsequently, biological evaluation revealed lowered cytotoxicity of all prepared amides (A1-A18) in relation to the MPA molecule, both for T-Jurkat and PBMC cell lines while A2 presented the most promising antiproliferative activity. Structure-activity relationship for differently substituted benzothiazoles and benzoxazoles was devised and indicated that amino group or substituent loci, electron-character of functionalities, and heteroatoms electronegativity might affected IC₅₀ and EC₅₀ values of newly synthesised derivers. These results confirm that the incorporation of aforementioned heterocyclic structural motifs into MPA core relevantly changed cytotoxicity and antiproliferative properties. Benzoxazole moiety in A2 turned out to be the most efficient in bioactivity alteration thus inclining further biological investigations in terms of different antiproliferative activities and structure-activity relationship determining. Benzothiazole MPA derivatives gave lower cytotoxicity. On the other hand, diminished toxicity against healthy cells may provide

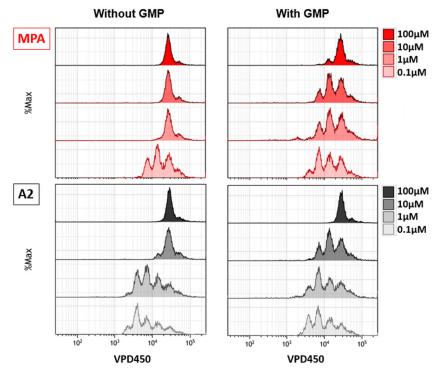


Figure 5. Representative antiproliferative activity of MPA and **A2**. VPD 450-labelled human PBMCs, in the presence of different concentrations (100; 10; 1 and 0.1μ M) of MPA (red) and **A2** (black) and stimulated (magnetic beads coated with anti-CD3 and anti-CD28 antibodies) were cultured with or without the addition of 50 μ M GMP for 72 h. Cell proliferation was analysed using flow cytometry.

active substance with improved therapeutic properties, including less side effects similarly to conjugate of MPA with quinic acid¹⁶. Moreover, investigations on both benzoxazole and benzothiazole MPA derivatives can lead to chemotherapeutics with anticancer potential^{32,33} causing them interesting structures for further exploring.

Experimental section

Computational studies

Ligands preparation

Prior to docking procedures, the potential inhibitors were prepared using Portable HyperChem 8.0.7 Release (Hypercube, Inc., Gainesville, FL^{48}). Each ligand was optimised using a MM + force field⁴⁹ and the Polak–Ribiere conjugate gradient algorithm (terminating at a gradient of 0.05 kcal mol⁻¹ Å⁻¹).

Protein preparation

The X-ray structures of the IMPDH II enzyme used for molecular modelling studies were taken from the Protein Databank (Protein Data Bank accession codes: 1JR1). The docking analysis was carried out after standard preparation procedures including removal of chain B of initial receptor structure, deletion of co-crystalised MPA structure and water molecules, and addition of hydrogen atoms and Gasteiger charges to each atom.

Molecular docking

Docking calculations of the optimised ligands to the prepared structure of human IMPDH II enzyme was carried out with Autodock Vina version 1.1.2 software (The Molecular Graphic Laboratory, The Scripps Research Institute, La Jolla, CA)⁵⁰. For all of docking studies, a grid box size was centred on the C β atom of

Chemistry

Urbana, IL)⁵¹.

General methods and materials

MPA, amines (1-18) used for A1-A18 synthesis, N,N-diisopropyle-(DIPEA), 1-hydroxybenzotriazole thylamine monohydrate $(HOBt \cdot H_2O),$ 2-(1*H*-benzotriazole-1-yl)-1,1,3,3-tetramethylaminium tetrafluoroborate (TBTU), 50% propylphosphonic acid anhydride in N,N-dimethylformamide (DMF; T3P) solution and pyridine (Py) were purchased from Sigma Aldrich, Apollo Scientific, AmBeed, Maybridge, Acros, Angene, Chemat, and Alfa Aesar and used as is. Organic solvents are commercially available on Sigma Aldrich, Fisher Scientific, Avantor, and Chempur stores. ACN, AcOEt, N,N-DMF, 1,4-dioxane (1,4-D), and Py were purified and dried upon standard procedures. Melting points (uncorrected) were measured using the Stuart SMP30 cryometer. ¹H and ¹³C NMR spectra were recorded on the Varian Unity Inova 500 spectrometer on the operating frequencies 500 MHz for ¹H NMR and 126 MHz for ¹³C NMR assays. Chemical shifts (δ) are given in parts per million (ppm) in relation to tetramethylsilane (TMS) with the residual solvent peak as internal standard (DMSO- d_6 : 2.50 ppm for ¹H NMR and 39.52 ppm for ¹³C NMR). Coupling constants (J) are shown in hertz (Hz). Residual solvent peaks, especially from ACN, dichloromethane, N,N-DMF, AcOEt, toluene, and water, may be observed and appear in typical values of chemical shifts. The description of individual peaks within ¹H and ¹³C NMR spectra was developed on the basis of the characteristic structural motifs present in MPA structure and the standard numbering brought under benzo[d]-1,3-azoles. Thin layer chromatography (TLC) analyses were carried

Ser276, and the size of the grid box was $25 \text{ Å} \times 25 \text{ Å} \times 25 \text{ Å}$.

Then, the best poses for a particular ligand were visually

inspected. Illustrations of the 3D model were generated using

VMD version 1.9 (University of Illinois at Urbana - Champaign,

out on Supelco silica gel TLC aluminium foils with fluorescence indicator 254 nm, whereas column chromatography was performed on silica gel 60 (230–450 Mesh, Alfa Aesar, Haverhill, MA). Visualisation of the chromatogram was achieved by UV light (254 nm) exposition or iodine staining.

HPLC analyses were performed using Agilent liquid chromatograph series 1290 (Agilent Technology, Waldbronn, Germany) consisting of binary pump G4220A, autosampler G4226A, thermostated column compartment G1316C, diode-array detector G1315C. The chromatographic system was controlled with Agilent MassHunter soft-ware B 06.01. The samples (2 µL) were injected onto a Poroshell EC-C18 2.7 µm (3.0 mm × 150 mm) column thermostated at 40 °C. The mobile phase flow rate was 0.4 mL·min⁻¹, and elution was performed using 0.1% (v/v) formic acid in water (solvent A) and MeCN/MeOH (1:1; v/v) (solvent B) in gradient mode: from 50% B to 100% B in 25 min. The UV signal was registered at 254 nm.

The accurate mass of the tested amides (**A1–A18**) and MPA was confirmed by electrospray ionisation quadrupole time-of-flight mass spectrometry ESI(-)-QTOF analysis using an Agilent 1290 LC system coupled to the Agilent QTOF mass spectrometer G6540 (Santa Clara, CA) operated in negative ionisation scan mode.

MPA characteristics present as follows: mp. 143–145 °C; ¹H NMR (DMSO–d₆) δ : 12.01 (s, 1H, –COOH), 9.40 (s, 1H, Ar–OH), 5.24 (s, 2H, Ar–CH₂O–), 5.13 (t, *J*=6.9 Hz, 1H, >C = CH–), 3.69 (s, 3H, Ar–OCH₃), 3.29 (d, *J*=6.8 Hz, 3H, Ar–CH₂–), 2.26 (t, *J*=7.4 Hz, 2H, –CH₂–), 2.16 (t, *J*=7.5 Hz, 2H, –CH₂), 2.08 (s, 3H, Ar–CH₃), 1.74 (s, 3H, –CH₃); ¹³C NMR (DMSO–d₆) δ 174.52 (–COOH), 170.58 (–COO–), 163.01 (C_{Ar}–OMe), 153.16 (C_{Ar}–OH), 146.27 (C_{Ar}–CH₂O–), 134.04 (>C = C(–CH₃)–), 123.13 (C_{Ar}–CH₂–CH=), 122.85 (>C = C(–CH₃)–), 116.40 (C_{Ar}–CH₃), 107.41 (C_{Ar}–COO–), 69.06 (Ar–CH₂O–), 61.08 (Ar–OCH₃), 34.57 (–CH₂–COOH), 32.91 (–CH₂–CH₂–), 22.87 (Ar–CH₂–), 16.46 (=C(–CH₃)–), 11.51 (Ar–CH₃); R_{T,HPLC} = 7.259 min, purity 100.00%; HRMS (*m*/*z*) calculated for C₁₇H₂₀O₆ [M — H]⁻: 319.1182; found: 319.1346.

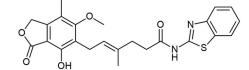
General methods for A1-A18 synthesis

We used two methods for syntheses of amide-like derivatives of MPA. Method A is intended for benzo-1,3-azoles aminated in the heteroaromatic region while Method B is dedicated for homoaromatic analogues. There are some differences in solvents used for carrying out these syntheses as well as some dissimilarities in applied reagents quantities. They are specified for particular cases.

Method A. In oven-dried, round bottom flask were placed 50 mg of MPA (0.156 mmol, 1.0 eg) and 27 mg of 97% 2-aminobenzo[d]thiazole (0.172 mmol, 1.1 eq). Afterwards, solids were dissolved in 1.5 mL of dry ACN and 0.05 mL of dry N,N-DMF. After being placed in an inert gas atmosphere (preferable argon) and cooled to 0°C with an ice bath, 55 μ L of *N*,*N*-DIPEA (d = 0.742 g/mL, 40.3 mg, 0.312 mmol, 2.0 eq) was added via syringe to the reaction system. Then, 24 mg of HOBt·H₂O (0.156 mmol, 1.0 eg) and 51 mg of 2-(1H-benzotriazole-1-yl)-1,1,3,3-TBTU (0.156 mmol, 1.0 eq) were added in one portion. Reaction mixture was allowed to heat to ambient temperature and was stirred for at least 48 h (preferably 72 h). Then, it was quenched with 0.4 mL of methanol, well mixed and filtered on the Hirsch funnel. White precipitate was thoroughly washed with cold solvents: ACN (ca. 2.0 mL), methanol (ca. 1.0 mL), and dichloromethane (ca. 1.0 mL) sequentially and well dried. White precipitate was then used without further purification while filtrate was discarded. A1 was obtained as off-white solid with 77% yield (54 mg).

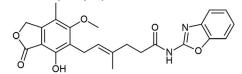
Method B. In oven-dried, round bottom flask was placed 125 mg of MPA (0.390 mmol, 1.0 eq) and 65 mg of 97% 6-aminobenzo[d]thiazole (0.429 mmol, 1.1 eq). Then, solids were dissolved in 2.0-3.0 mL of dry AcOEt and the reaction system was cooled down to 0°C with an ice bath. Under an inert gas atmosphere, 63 μ L of dry Py (d = 0.982 g/mL, 61.7 mg, 0.780 mmol, 2.0 eg) was added via syringe to the reaction mixture. Afterwards, 342 µL of 50% solution of propylphosphonic acid anhydride in N,N-DMF (T3P) (d = 1.09 g/mL, 186.2 mg, 0.585 mmol, 1.5 eg) was added dropwise in 30 min. Then it was allowed to stir for at least 48 h (preferably 72 h) at room temperature. This mixture was then diluted with 5 mL of fresh AcOEt, extracted with 1 M hydrochloric acid solution, saturated sodium carbonate solution and then washed with brine. Organic phase was dried over anhydrous magnesium sulphate, filtered off, concentrated under reduced pressure and then purified over column chromatography in toluene/AcOEt system (2/1, v/v, $R_F = 0.13$). A4 was obtained as a fluffy, ecru solid with 55% yield (97 mg).

N-(benzo[d]thiazol-2-yl) mycophenolate (A1).



Method A - 2.0 eq DIPEA, afforded the product as white solid with 77% yield; mp. 205–210 °C; ¹H NMR (DMSO–d₆) δ 12.23 (s, 1H, -NH-), 9.35 (s, 1H, Ar-OH), 7.94 (d, J=7.8 Hz, 1H, CH_{HetAr}), 7.71 (d, J = 8.0 Hz, 1H, CH_{HetAr}), 7.43 (t, J = 7.6 Hz, 1H, CH_{HetAr}), 7.30 (t, J = 7.5 Hz, 1H, CH_{HetAr}), 5.18 (t, J = 6.4 Hz, 1H, $>C = CH_{-}$), 5.04 (s, 2H, Ar-CH₂O-), 3.65 (s, 3H, Ar-OCH₃), 3.28 (d, J=6.7 Hz, 2H, Ar-CH₂-), 2.56 (t, J = 7.4 Hz, 2H, -CH₂-), 2.29 (t, J = 7.2 Hz, 2H, -CH₂-), 1.96 (s, 3H, Ar-CH₃), 1.78 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 172.19 (–CONH–), 170.56 (–COO–), 162.83 (C_{Ar}–OMe), 158.25 (C(2)_{HetAr}), 153.06 (C_{Ar}-OH), 148.91 (C(3a)_{HetAr}), 146.08 $(C_{Ar}-CH_2\overline{O}-)$, 133.62 (>C = $\overline{C}(-CH_3)-$), 131.87 ($\overline{C}(7a)_{HetAr}$), 126.45 (C(5)_{HetAr}), 123.83 (C(6)_{HetAr}), 123.64 (C_{Ar}-CH₂-CH=), 122.65 $(>C = C(-CH_3)-)$, 122.02 (C(7)_{HetAr}), 120.85 (C(4)_{HetAr}), 116.23 (C_{Ar}-CH₃), 107.26 (C_{Ar}-COO-), 68.88 (Ar-CH₂O-), 60.97 (Ar-OCH₃), 34.64 (-CH₂-CONH-), 34.33 (-CH₂-CH₂-), 22.88 (Ar-CH₂-), 16.50 $(=C(-CH_3)-)$, 11.40 (Ar-CH₃); $R_{T,HPLC} = 12.253$ min, purity 98.95%; HRMS (m/z) calculated for $C_{24}H_{24}N_2O_5S$ $[M - H]^-$: 451.1328; found: 451.1572.

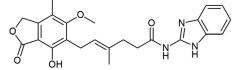
N-(benzo[d]oxazol-2-yl) mycophenolate (A2).



Method A – 1.0 eq DIPEA, washing step limited to cold ACN usage, afforded the product as white solid with 72% yield; mp. 175–177 °C; ¹H NMR (DMSO– d_6) δ 11.57 (bs, 1H, –NH–), 9.40 (bs, 1H, Ar–OH), 7.64–7.49 (m, 2H, CH_{HetAr}), 7.28 (m, 2H, CH_{HetAr}), 5.22–5.13 (m, 3H, >C = CH– and Ar–CH₂O–), 3.67 (s, 3H, Ar–OCH₃), 3.30 (d, *J* = 6.8 Hz, 2H, Ar–CH₂–), 2.58 (m, 2H, –CH₂–), 2.27 (t, *J* = 7.5 Hz, 2H, –CH₂–), 2.02 (s, 3H, Ar–CH₃), 1.78 (s, 3H, –CH₃); ¹³C NMR (DMSO– d_6) δ 170.56 (–COO–), 162.95 (C_{Ar}–OMe), 155.49 (C(2)_{HetAr}), 153.12 (C_{Ar}–OH), 147.96 (C(7a)_{HetAr}), 146.20 (C_{Ar}–CH₂O–), 141.10 (C(3a)_{HetAr}), 133.84 (>C = C(–CH₃)–), 124.92 (C(6)_{HetAr}), 123.86 (C(5)_{HetAr}), 123.48 (C_{Ar}–CH₂–CH=), 122.77 (>C = C(–CH₃)–), 118.55 (C(4)_{HetAr}), 116.33 (C_{Ar}–CH₃), 110.38 (C(7)_{HetAr}), 107.36 (C_{Ar}–COO–), 68.99 (Ar–CH₂O–), 61.03 (Ar–OCH₃),

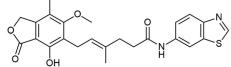
35.08 (-CH₂-CONH-), 34.50 (-CH₂-CH₂-), 22.89 (Ar-CH₂-), 16.57 (=C(-CH₃)-), 11.45 (Ar-CH₃); $R_{T,HPLC} = 9.474$ min, purity 98.96%; HRMS (*m/z*) calculated for $C_{24}H_{24}N_2O_6$ [M — H]⁻: 435.1556; found: 435.1793.

N-(1H-benzo[d]imidazol-2-yl) mycophenolate (A3).



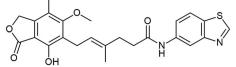
Method A - 1.5 eq DIPEA, 1.5 eq TBTU, without HOBt·H₂O, with additional washing step with cold DMF, afforded the product as fluffy, white solid with 43% yield; mp. 253-255 °C (with decomposition); ^1H NMR (DMSO–d_6) δ 11.90 (bs, 1H, –NH–), 11.39 (bs, 1H, NH_{HetAr}), 9.44 (bs, 1H, Ar-OH), 7.45 - 7.36 (m, 2H, CH_{HetAr}), 7.06 (dd, J = 5.8, 3.1 Hz, 2H, CH_{HetAr}), 5.19 (t, J = 6.6 Hz, 1H, >C = CH-), 5.12 (s, 2H, Ar-CH₂-), 3.63 (s, 3H, Ar-OCH₃), 3.29 (d, J = 6.8 Hz, 2H, Ar-CH₂-), 2.52 (d, J = 6.5 Hz, 2H, -CH₂-), 2.30 (t, J = 7.4 Hz, 2H, $-CH_2-$), 1.97 (s, 3H, Ar $-CH_3$), 1.79 (s, 3H, $-CH_3$); ¹³C NMR (DMSO-d₆) δ 183.06 (C(2)_{HetAr}), 172.40 (–CONH–), 170.60 (-<u>C</u>OO-), 162.87 (C_{Ar}-OMe), 153.06 (C_{Ar}-OH), 146.98 (C_{HetAr}), $(C_{Ar}-CH_2\overline{O}-),$ 146.16 133.79 $(>C = C(-CH_3)-),$ 123.62 $(C_{Ar}-CH_2-CH=)$, 122.71 (>C = C(-CH_3)-), 121.29 (C_{HetAr}), 116.29 (C_{Ar}-CH₃), 107.33 (C_{Ar}-COO-), 68.96 (Ar-CH₂O-), 60.97 (Ar-OCH₃), 34.86 (-CH₂-CONH-), 34.56 (-CH₂-CH₂-), 22.88 (Ar-CH₂-), 16.48 $(=C(-CH_3)-)$, 11.40 (Ar-CH₃); $R_{T,HPLC} = 5.617 \text{ min}$, purity 99.50%; HRMS (*m/z*) calculated for $C_{24}H_{25}N_3O_5$ [M – H]⁻: 434.1716; found: 434.1953.

N-(benzo[d]thiazol-6-yl) mycophenolate (A4).



Method B afforded the product as fluffy, creamy solid with 55% yield; mp. 175–177 °C; ¹H NMR (DMSO–d₆) δ 10.11 (s, 1H, –NH–), 9.42 (bs, 1H, C(2)H_{HetAr}), 9.25 (s, 1H, Ar–OH), 8.47 (d, J = 1.7 Hz, 1H, C(7)H_{HetAr}), 7.95 (d, J = 8.8 Hz, 1H, CH_{HetAr}), 7.48 (dd, J = 8.8, 1.9 Hz, 1H, CH_{HetAr}), 5.23 – 5.15 (m, 3H, $>\!C\!=\!CH\!-$ and Ar–CH_2O–), 3.64 (s, 3H, Ar-OCH₃), 3.30 (d, J=6.8 Hz, 2H, Ar-CH₂-), 2.41 (t, J=7.6 Hz, 2H, -CH₂-), 2.28 (t, J=7.5 Hz, 2H, -CH₂-), 2.00 (s, 3H, Ar-CH₃), 1.79 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 171.46 (-CONH-), 170.58 (-COO-), 162.89 (C_{Ar}-OMe), 154.92 (C(2)_{HetAr}), 153.14 (C_{Ar}-OH), 149.31 (C(3a)_{HetAr}), 146.13 (C_{Ar}-CH₂O-), 137.42 (C(6)_{HetAr}), 134.58 $(>C = C(-CH_3)-)$, 134.11 (C(7a)_{HetAr}), 123.27 (C_{Ar}-CH₂-CH=), 123.24 60.99 (Ar-OCH₃), 35.66 (-CH₂-CONH-), 35.21 (-CH₂-CH₂-), 22.91 $(Ar-CH_2-)$, 16.64 (=C(-CH_3)-), 11.43 (Ar-CH_3); $R_{T,HPLC} = 8.638 \text{ min}$, purity 100.00%; HRMS (m/z) calculated for C₂₄H₂₄N₂O₅S $[M - H]^-$: 451.1328; found: 451.1572.

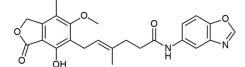
N-(benzo[d]thiazol-5-yl) mycophenolate (A5).



Method B – column chromatography carried out in toluene/AcOEt mixture (2/1, v/v, $R_F = 0.18$), afforded the product as fluffy, white

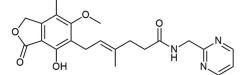
solid with 67% yield; mp. 114–117 °C; ¹H NMR (DMSO–d₆) δ 10.08 (s, 1H, -NH-), 9.26 (m, 2H, Ar-OH and C(2)H_{HetAr}), 8.38 (d, J = 1.8 Hz, 1H, CH_{HetAr}), 8.01 (d, J = 8.7 Hz, 1H, CH_{HetAr}), 7.50 (dd, J = 8.7, 1.9 Hz, 1H, CH_{HetAr}), 5.22 – 5.16 (m, 3H, >C = CH- and Ar-CH₂O-), 3.64 (s, 3H, Ar–OCH₃), 3.30 (d, J = 6.9 Hz, 2H, Ar–CH₂–), 2.41 (dd, J = 8.9, 6.4 Hz, 2H, -CH₂-), 2.28 (t, J = 7.6 Hz, 2H, -CH₂-), 2.00 (s, 3H, Ar-CH₃), 1.79 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 171.44 (-CONH-), 170.60 (-COO-), 162.89 (C_{Ar}–OMe), 157.24 (C(2)_{HetAr}), 153.98 (C(3a)_{HetAr}), 153.12 (C_{Ar}–OH), 146.15 (C_{Ar}–CH₂O–), 138.29 $(C(5)_{HetAr}),$ 134.14 $(>C = C(-CH_3)-)$, 127.98 (C(7a)_{HetAr}), 123.26 (C_{Ar}-CH₂-CH=), 122.82 $(C(6)_{HetAr})$, 122.57 (>C = C(-CH₃)-), 118.33 (C(7)_{HetAr}), 116.32 (C_{Ar}-CH₃), 112.98 (C(4)_{HetAr}), 107.34 (C_{Ar}-COO-), 69.02 (Ar-CH₂O-), 60.98 (Ar-OCH₃), 35.70 (-CH₂-CONH-), 35.17 (-CH₂-CH₂-), 22.90 (Ar-CH₂-), 16.66 (=C(-CH₃)-), 11.41 (Ar-CH₃); R_{T,HPLC} = 8.651 min, purity 100.00%; HRMS (*m/z*) calculated for C₂₄H₂₄N₂O₅S [M — H]⁻: 451.1328; found: 451.1573.

N-(benzo[d]oxazol-5-yl) mycophenolate (A6).



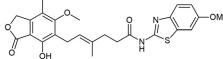
Method B - column chromatography carried out in toluene/acetone mixture (3/1, v/v, $R_F = 0.34$), afforded the product as beige solid with 53% yield; mp. 84–87 °C; ¹H NMR (DMSO–d₆) δ 10.01 (s, 1H, -NH-), 9.37 (s, 1H, Ar-OH), 8.69 (s, 1H, C(2)H_{HetAr}), 8.05 (d, J = 1.9 Hz, 1H, C(4)H_{HetAr}), 7.64 (d, J = 8.8 Hz, 1H, CH_{HetAr}), 7.43 (dd, J = 8.8, 2.0 Hz, 1H, CH_{HetAr}), 5.19 (m, 3H, >C = CH- and Ar-CH₂O-), 3.64 (s, 3H, Ar-OCH₃), 3.30 (d, J=6.8 Hz, 2H, Ar-CH₂-), 2.45 - 2.35 (m, 2H, -CH₂-), 2.28 (dd, J=13.4, 5.3 Hz, 2H, -CH₂-), 2.01 (s, 3H, Ar-CH₃), 1.79 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 171.21 (-CONH-), 170.59 (-COO-), 162.91 (C_{Ar}-OMe), 155.22 (C(2)_{HetAr}), 153.13 (C_{Ar}-OH), 146.16 (C_{Ar}-CH₂O-), 145.68 (C(3a)_{HetAr}), 140.19 $(C(7a)_{HerAr})$, 136.73 $(C(5)_{HetAr})$, 134.16 $(>C = C(-CH_3)-)$, 123.23 $(C_{Ar}-CH_2-CH=)$, 122.83 (>C = C(-CH_3)-), 118.00 (C(6)_{HetAr}), 116.33 (C_{Ar}-CH₃), 111.18 (C_{HetAr}), 110.54 (C_{HetAr}), 107.35 (C_{Ar}-COO-), 69.03 (Ar-CH₂O-), 60.98 (Ar-OCH₃), 35.62 (-CH₂-CONH-), 35.20 (-CH₂-CH₂-), 22.90 (Ar-CH₂-), 16.64 (=C(-CH₃)-), 11.43 (Ar-CH₃); $R_{T,HPLC} = 7.866 \text{ min}$, purity 95.79%; HRMS (*m/z*) calculated for C₂₄H₂₄N₂O₆ [M — H]⁻: 435.1556; found: 435.1795.

N-[1-(pirymidin-2-yl)methyl] mycophenolate (A7).



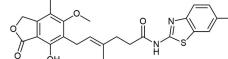
Method B - column chromatography carried out in toluene/acetone mixture (2/1, v/v, $R_F = 0.13$), afforded the product as white solid with 61% yield; mp. 107–110 °C; ¹H NMR (DMSO–d₆) δ 9.41 (s, 1H, Ar-OH), 8.73 (d, J=4.9 Hz, 2H, CH_{HetAr}), 8.33 (t, J=5.8 Hz, 1H, -NH-), 7.37 (t, J = 4.9 Hz, 1H, CH_{HetAr}), 5.24 (s, 2H, Ar-CH₂O-), 5.14 (t, J = 6.9 Hz, 1H, >C = CH-), 4.40 (d, J = 5.8 Hz, 2H, -CH₂-HetAr), 3.69 (s, 3H, Ar-OCH₃), 3.30 (d, J=8.3 Hz, 2H, Ar–CH₂–), 2.28 - 2.21 (m, 2H, –CH₂–), 2.17 (dd, J = 8.8, 5.6 Hz, 2H, -CH₂-), 2.08 (s, 3H, Ar-CH₃), 1.75 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 172.34 (-CONH-), 170.59 (-COO-), 167.48 (C(2)_{HetAr}), 163.00 (C_{Ar}–OMe), 157.72 (C(4)_{HetAr} and C(6)_{HetAr}), 153.17 (C_{Ar}–OH), 146.23 (C_{Ar}-CH₂O-), 134.52 $(>C = C(-CH_3)-),$ 123.01 (C_{Ar}-CH₂-CH=), 122.94 (>C = C(-CH₃)-), 120.17 (C(5)_{HetAr}), 116.39 (<u>C</u>_{Ar}-CH₃), 107.41 (<u>C</u>_{Ar}-COO-), 69.06 (Ar-<u>C</u>H₂O-), 61.07 (Ar-OCH₃),

N-(6-methoxybenzo[d]thiazol-2-yl) mycophenolate (A8).



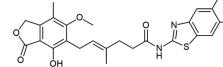
Method A - 2.0 eq DIPEA, afforded the product as white solid with 100% yield; mp. 246-248 °C (with decomposition); ¹H NMR (DMSO-d₆) δ 12.10 (s, 1H, -NH-), 9.35 (s, 1H, Ar-OH), 7.60 (d, J = 8.8 Hz, 1H, CH_{HetAr}), 7.53 (d, J = 2.4 Hz, 1H, CH_{HetAr}), 7.02 (dd, J = 8.8, 2.5 Hz, 1H, CH_{HetAr}), 5.17 (t, J = 6.7 Hz, 1H, $>C = CH_{-}$), 5.08 (s, 2H, Ar-CH₂O-), 3.81 (s, 3H, HetAr-OCH₃), 3.65 (s, 3H, Ar-OCH₃), 3.28 (d, J = 6.8 Hz, 2H, Ar-CH₂-), 2.58 - 2.52 (m, 2H, -CH₂-), 2.28 (t, J=7.4 Hz, 2H, -CH₂-), 1.98 (s, 3H, Ar-CH₃), 1.78 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 171.89 (-CONH-), 170.61 (-COO-), 162.94 (C_{Ar}-OMe), 156.51 (C(2)_{HetAr}), 156.28 (C(6)_{HetAr}), 153.11 (C_{Ar}-OH), 146.06 (C_{Ar} -CH₂O-), 143.07 (C(3a)_{HetAr}), 133.72 (>C = C(-CH₃)-), 133.21 ($\overline{C}(7a)_{HetAr}$), 123.61 ($C_{Ar}-CH_2-CH=$), 122.65 (>C = C(-CH_3)-), 121.44 (C(4)_{HetAr}), 116.30 (C_{Ar}-CH₃), 115.19 (C(5)_{HetAr}), 107.29 (C_{Ar}-COO-), 105.13 (C(7)_{HetAr}), 68.95 (Ar-CH₂O-), 61.01 (Ar-OCH₃), 56.12 (HetAr–OCH₃), 34.69 (–CH₂–CONH–), 34.35 (–CH₂–CH₂–), 22.89 (Ar-CH₂-), 16.50 (=C($-CH_3$)-), 11.42 (Ar-CH₃); $R_{T,HPLC}$ = 12.073 min, purity 99.43%; HRMS (m/z) calculated for C₂₅H₂₆N₂O₆S [M — H]⁻: 481.1433; found: 481.1692.

N-(6-methylbenzo[d]thiazol-2-yl) mycophenolate (A9).



Method A - 1.0 eq DIPEA, afforded the product as white solid with 81% yield; mp. 257–260 °C (with decomposition); ¹H NMR (DMSO-d_6) δ 12.15 (s, 1H, -NH-), 9.38 (s, 1H, Ar-OH), 7.72 (s, 1H, C(7)H_{HetAr}), 7.59 (d, J = 8.2 Hz, 1H, CH_{HetAr}), 7.24 (d, J = 8.2 Hz, 1H, CH_{HetAr}), 5.17 (t, J = 6.7 Hz, 1H, $>C = CH_{-}$), 5.06 (s, 2H, Ar- $CH_{2}O_{-}$), 3.67 (s, 3H, Ar-OCH₃), 3.28 (d, J=6.9 Hz, 2H, Ar-CH₂-), 2.53 (m, 2H, -CH₂-), 2.41 (s, 3H, HetAr-CH₃), 2.27 (t, J = 7.4 Hz, 2H), 1.97 (s, 3H, Ar–CH₃), 1.76 (s, 3H, –CH₃); ¹³C NMR (DMSO–d₆) δ 172.03 (-CONH-), 170.67 (-COO-), 163.03 (C_{Ar}-OMe), 157.42 (C(2)_{HetAr}), 153.14 (C_{Ar}-OH), 146.97 (C(3a)_{HetAr}), 146.02 (C_{Ar}-CH₂O-), 133.73 $(>C = C(-CH_3)-)$, 133.25 $(C(7a)_{HetAr})$, 132.12 $(C(6)_{HetAr})$, 127.71 $(C(5)_{HetAr})$, 123.64 $(C_{Ar}-CH_2-CH=)$, 122.63 $(>C=C(-CH_3)-)$, 121.55 (C_{HetAr}), 120.50 (C_{HetAr}), 116.33 (C_{Ar}-CH₃), 107.30 (C_{Ar}-COO-), 68.99 (Ār-CH₂O-), 61.03 (Ar-OCH₃), 34.69 (-CH₂-CONH-), 34.46 (-CH₂-CH₂-), 22.89 (Ar-CH₂-), 21.39 (HetAr-CH₃), 16.48 $(=C(-CH_3)-)$, 11.41 (Ar-CH₃); $R_{T,HPLC} = 13.755$ min, purity 98.99%; HRMS (m/z) calculated for C₂₅H₂₆N₂O₅S $[M - H]^-$: 465.1484; found: 465.1721.

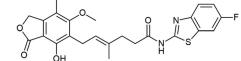
N-(5,6-dimethylbenzo[d]thiazol-2-yl) mycophenolate (A10).



Method A – 1.0 eq DIPEA, afforded the product as white solid with 77% yield; mp. 246–247 $^\circ\text{C}$ (with decomposition); ^1H NMR (DMSO–d₆) δ 12.12 (s, 1H, –NH–), 9.36 (s, 1H, Ar–OH), 7.66 (s, 1H,

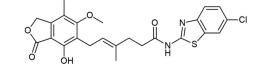
CH_{HetAr}), 7.50 (s, 1H, CH_{HetAr}), 5.17 (t, J = 6.6 Hz, 1H, >C = CH-), 5.08 (s, 2H, Ar–CH₂O–), 3.64 (s, 3H, Ar–OCH₃), 3.28 (d, J = 6.7 Hz, 2H, Ar–CH₂–), 2.57 – 2.51 (m, 2H, –CH₂–), 2.33 (m, 8H, –CH₂–, HetAr(5)–CH₃, and HetAr(6)–CH₃), 1.96 (s, 3H, Ar–CH₃), 1.78 (s, 3H, –CH₃); ¹³C NMR (DMSO–d₆) δ 171.93 (–CONH–), 170.57 (–COO–), 162.86 (C_{Ar}–OMe), 157.33 (C(2)_{HetAr}), 153.08 (C_{Ar}–OH), 147.48 (C(3a)_{HetAr}), 146.13 (C_{Ar}–CH₂O–), 135.06 (C(7a)_{HetAr}), 133.70 ($>C = C(-CH_3)$ –), 132.64 (C(6)_{HetAr}), 129.22 (C(5)_{HetAr}), 123.59 (C_{Ar}–CH₂–CH=), 122.67 ($>C = C(-CH_3)$ –), 121.77 (C(7)_{HetAr}), 121.24 (C(4)_{HetAr}), 116.26 (C_{Ar}–CH₃), 107.30 (C_{Ar}–COO–), 68.90 (Ar–CH₂O–), 60.99 (Ar–OCH₃), 34.68 (–CH₂–CONH–), 34.32 (–CH₂–CH₂–C), 22.88 (Ar–CH₂–), 20.17 (HetAr–CH₃), 20.01 (HetAr–CH₃), 16.50 (=C(–CH₃)–), 11.42 (Ar–CH₃); R_{T,HPLC} = 14.909 min, purity 97.49%; HRMS (*m*/z) calculated for C₂₆H₂₈N₂O₅S [M – H]⁻: 479.1641; found: 479.1893.

N-(6-fluorobenzo[d]thiazol-2-yl) mycophenolate (A11).



Method A - 2.0 eq DIPEA, afforded the product as white solid with 71% yield; mp. 232–236°C (with decomposition); ¹H NMR (DMSO-d₆) δ 12.26 (s, 1H, -NH-), 9.35 (s, 1H, Ar-OH), 7.86 (dd, J=8.7, 2.5 Hz, 1H, CH_{HetAr}), 7.71 (dd, J=8.8, 4.8 Hz, 1H, CH_{HetAr}), 7.28 (td, J=9.1, 2.6 Hz, 1H, CH_{HetAr}), 5.17 (t, J=6.7 Hz, 1H, >C = CH-), 5.08 (s, 2H, Ar-CH₂O-), 3.65 (s, 3H, Ar-OCH₃), 3.28 (d, J = 6.8 Hz, 2H, Ar-CH₂-,), 2.55 (t, J = 7.5 Hz, 2H, -CH₂-), 2.29 (t, J = 7.4 Hz, 2H, $-CH_2-$), 1.98 (s, 3H, Ar $-CH_3$), 1.78 (s, 3H, $-CH_3$); ¹³C NMR (DMSO-d₆) δ 172.26 (-CONH-), 170.66 (-COO-), 163.04 $(C_{Ar}-OMe)$, 160.02 $(C(2)_{HetAr})$, 158.20 $(d, J = 21.3 \text{ Hz}, C(6)_{HetAr})$, 153.14 (C_{Ar}-OH), 145.98 (C_{Ar}-CH₂O-), 145.71 (C(3a)_{HetAr}), 133.70 $(>C = C(-CH_3)-)$, 133.20 (d, J = 11.1 Hz, $C(7a)_{\text{HetAr}}$), 123.65 $(C_{Ar}-CH_2-CH=)$, 122.63 (>C = C(-CH_3)-), 121.90 (d, J = 9.2 Hz, $\overline{C(4)}_{HetAr}$), 116.34 (C_{Ar}-CH₃), 114.45 (d, J = 24.5 Hz, C(5)_{HetAr})), 108.37 (d, J = 26.8 Hz, C(7)_{HetAr}), 107.28 (C_{Ar}-COO-), 69.02 (Ar-CH₂O-), 61.02 (Ar-OCH₃), 34.63 (-CH₂-CONH-), 34.43 (-CH₂-CH₂-), 22.89 (Ar-CH₂-), 16.47 (=C(-CH₃)-), 11.40 (Ar-CH₃); $R_{T,HPLC} = 12.970 \text{ min}$, purity 99.58%; HRMS (*m/z*) calculated for C₂₄H₂₃FN₂O₅S [M — H]⁻: 469.1233; found: 469.1488.

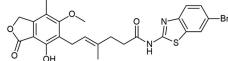
N-(6-chlorobenzo[d]thiazol-2-yl) mycophenolate (A12).



Method A – 1.0 eq DIPEA, afforded the product as white solid with 78% yield; mp. 261–264 °C (with decomposition); ¹H NMR (DMSO–d₆) δ 12.33 (s, 1H, –NH–), 9.38 (s, 1H, Ar–OH), 8.09 (s, 1H, C(7)H_{HetAr}), 7.70 (d, *J* = 8.6 Hz, 1H, CH_{HetAr}), 7.45 (d, *J* = 8.5 Hz, 1H, CH_{HetAr}), 5.17 (t, *J* = 6.2 Hz, 1H, >C = CH–), 5.08 (s, 2H, Ar–CH₂O–), 3.67 (s, 3H, Ar–OCH₃), 3.28 (d, *J* = 6.6 Hz, 2H, Ar–CH₂–), 2.56 (t, *J* = 7.4 Hz, 2H, –CH₂–), 2.29 (t, *J* = 7.1 Hz, 2H, –CH₂–), 1.97 (s, 3H, Ar–CH₃), 1.78 (s, 3H,–CH₃); ¹³C NMR (DMSO–d₆) δ 172.40 (–CONH–), 170.61 (–COO–), 162.96 (C_{Ar}–OMe), 159.12 (C(2)_{HetAr}), 153.12 (C_{Ar}–OH), 147.86 (C(3a)_{HetAr}), 146.02 (C_{Ar}–CH₂O–), 133.64 (>C = C(–CH₃)–), 122.64 (>C = C(–CH₃)–), 122.04 (C_{HetAr}), 121.69 (C_{Ar}–CH₂–CH=), 122.64 (>C = C(–CH₃)–), 122.04 (C_{HetAr}), 121.69 (C_{HetAr}), 116.30 (C_{Ar}–CH₃), 107.28 (C_{Ar}–COO–), 68.96 (Ar–CH₂O–), 61.01 (Ar–OCH₃), 34.62 (–CH₂–CONH–), 34.41 (–CH₂–CH₂–), 22.89 (Ar–CH₂–), 16.48 (=C(–CH₃)–), 11.41 (Ar–CH₃); R_{T,HPLC} = 14.845 min,

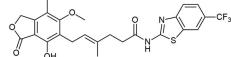
purity 97.55%; HRMS (*m/z*) calculated for $C_{24}H_{23}CIN_2O_5S$ [M — H]⁻: 485.0938; found: 485.1186.

N-(6-bromobenzo[d]thiazol-2-yl) mycophenolate (A13).



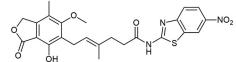
Method A - 1.0 eq DIPEA, additional 0.15 mL of dry 1,4-D was added to enhance solubility, afforded the product as creamy solid with 80% yield; mp. 265-267 °C (with decomposition); ¹H NMR (DMSO-d₆) δ 12.33 (s, 1H, -NH-), 9.37 (s, 1H, Ar-OH), 8.22 (d, J = 1.2 Hz, 1H, CH_{HetAr}), 7.67 (d, J = 8.6 Hz, 1H, CH_{HetAr}), 7.57 (dd, J = 8.6, 1.4 Hz, 1H, CH_{HetAr}), 5.17 (t, J = 6.6 Hz, 1H, >C = CH-), 5.08 (s, 2H, Ar-CH₂O-), 3.64 (s, 3H, Ar-OCH₃), 3.28 (d, J=6.8 Hz, 2H, Ar–CH₂–), 2.56 (t, J = 7.4 Hz, 2H, –CH₂–), 2.29 (t, J = 7.4 Hz, 2H, -CH₂-), 1.97 (s, 3H, Ar-CH₃), 1.76 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 172.41 (-CONH-), 170.65 (-COO-), 163.02 (C_{Ar}-OMe), 159.10 (C(2)_{HetAr}), 153.14 (C_{Ar}-OH), 148.18 (C(3a)_{HetAr}), 145.99 $(C_{Ar}-CH_2O-)$, 134.18 $(C(7a)_{HetAr})$, 133.66 $(>C=C(-CH_3)-)$, 129.45 (C(7a)_{HetAr}), 124.50 (C_{HetAr}), 123.68 (C_{Ar}-CH₂-CH=), 122.62 $(>C = C(-CH_3)-)$, 122.45 (C_{HetAr}) , 116.33 $(C_{Ar}-CH_3)$, 115.73 (C(6)_{HetAr}), 107.29 (C_{Ar}-COO-), 68.99 (Ar-CH₂O-), 61.02 (Ar-OCH₃), 34.63 (-CH₂-CONH-), 34.46 (-CH₂-CH₂-), 22.89 (Ar-CH₂-), 16.47 $(=C(-CH_3)-)$, 11.40 (Ar-CH₃); $R_{T,HPLC} = 15.319$ min, purity 97.88%; HRMS (m/z) calculated for C₂₄H₂₃BrN₂O₅S $[M - H]^-$: 529.0433, 531.0415; found: 529.0703, 531.0687.

N-[6-(trifluoromethyl)benzo[d]thiazol-2-yl] mycophenolate (A14).



Method A - 2.0 eq DIPEA, afforded the product as white solid with 65% yield; mp. 261-264 °C (with decomposition); ¹H NMR (DMSO-d_6) δ 12.49 (s, 1H, -NH), 9.34 (s, 1H, Ar-OH), 8.47 (s, 1H, C(7)H_{HetAr}), 7.87 (d, J=8.5 Hz, 1H, CH_{HetAr}), 7.74 (d, J=8.5 Hz, 1H, CH_{HetAr}), 5.18 (t, J = 6.8 Hz, 1H, >C = CH-), 5.05 (s, 2H, Ar- CH_2O -), 3.64 (s, 3H, Ar-OCH₃), 3.28 (d, J = 6.9 Hz, 2H, Ar-CH₂-), 2.59 (t, J=7.5 Hz, 2H, -CH₂-), 2.37 - 2.24 (m, 2H, -CH₂-), 1.96 (s, 3H, Ar-CH₃), 1.79 (s, 3H, -CH₃); ¹³C NMR (DMSO-d₆) δ 172.66 (-CONH-), 170.59 (-COO-), 162.96 (C_{Ar}-OMe), 161.47 (C(2)_{HetAr}), 153.11 (C_{Ar}-OH), 151.74 (C(3a)_{HetAr}), 145.99 (C_{Ar}-CH₂O-), 133.60 $(>C = C(-CH_3)-)$, 132.50 $(\overline{C}(7a)_{HetAr})$, 126.14 (\overline{C}_{HetAr}) , 124.09 (d_r) J = 31.9 Hz, HetAr-CF₃) 123.70 (C_{Ar}-CH₂-CH=), 123.24 (d, $J = 3.4 \text{ Hz}, C_{\text{HetAr}}$, $1\overline{22.63}$ (>C = C(-CH₃)-), 121.24 (C_{HetAr}), 120.14 (d, J = 4.1 Hz, C(4)_{HetAr}), 116.28 (C_{Ar}-CH₃), 107.27 (C_{Ar}-COO-), 68.91 (Ar-CH₂O-), 61.00 (Ar-OCH₃), 34.57 (-CH₂-CONH-), 34.44 $(-CH_2^--CH_2^-)$, 22.89 (Ar- CH_2^-), 16.48 (= $C(-CH_3^-)$), 11.37 (Ar- CH_3); $R_{T,HPLC} = 15.435 \text{ min}$, purity 97.28%; HRMS (*m/z*) calculated for $C_{25}H_{23}F_{3}N_{2}O_{5}S$ [M — H]⁻: 519.1202; found: 519.1484.

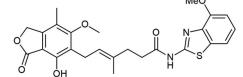
N-(6-nitrobenzo[d]thiazol-2-yl) mycophenolate (A15).



Method A – 1.0 eq DIPEA, reaction was carried out in 0.6 mL of dry DMF and 1.0 mL of dry ACN, afforded the product as beige solid with 42% yield; mp. 278–280 °C (with decomposition); ¹H NMR (DMSO-d₆) δ 12.66 (s, 1H, –NH–), 9.37 (s, 1H, Ar–OH), 9.03 (d,

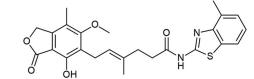
 $\begin{array}{l} J=2.0\,\text{Hz},\ 1\text{H},\ C\text{H}_{\text{HetAr}}\text{I},\ 8.28\ (dd,\ J=8.0,\ 4.0\,\text{Hz},\ 1\text{H},\ C\text{H}_{\text{HetAr}}\text{I},\ 7.86\ (d,\ J=8.9\,\text{Hz},\ 1\text{H},\ C\text{H}_{\text{HetAr}}\text{I},\ 5.18\ (t,\ J=6.6\,\text{Hz},\ 1\text{H},\ >C=C\text{H}),\ 5.08\ (s,\ 2\text{H},\ Ar-C\text{H}_2\text{O}),\ 3.65\ (s,\ 3\text{H},\ Ar-OC\text{H}_3\text{J},\ 3.28\ (d,\ J=6.7\,\text{Hz},\ 2\text{H},\ Ar-C\text{H}_2\text{O}),\ 2.61\ (t,\ J=7.4\,\text{Hz},\ 2\text{H},\ -C\text{H}_2\text{-}),\ 2.31\ (t,\ J=7.3\,\text{Hz},\ 2\text{H},\ -C\text{H}_2\text{-}),\ 2.61\ (t,\ J=7.4\,\text{Hz},\ 2\text{H},\ -C\text{H}_2\text{-}),\ 2.31\ (t,\ J=7.3\,\text{Hz},\ 2\text{H},\ -C\text{H}_2\text{-}),\ 1.97\ (s,\ 3\text{H},\ Ar-C\text{H}_3\text{J}),\ 1.80\ (s,\ 3\text{H},\ -C\text{H}_3\text{J});\ ^{13}\text{C}\ NMR\ (DMSO-d_6)\ \delta\ 172.96\ (-CONH-),\ 170.57\ (-COO-),\ 163.82\ (C(2)_{\text{HetAr}}\text{J}),\ 162.98\ (C_{Ar}-OMe),\ 153.92\ (C(6)_{\text{HetAr}}\text{J}),\ 153.12\ (C_{Ar}-O\text{H}),\ 146.02\ (C_{Ar}-C\text{H}_2\text{O}-),\ 143.41\ (C(3a)_{\text{HetAr}}\text{J}),\ 133.58\ (>C=C(-C\text{H}_3)-),\ 132.66\ (\overline{C}(7a)_{\text{HetAr}}\text{J},\ 123.73\ (C_{Ar}-C\text{H}_2-C\text{H}=),\ 122.63\ (>C=C(-C\text{H}_3)-),\ 122.12\ (\overline{C}(5)_{\text{HetAr}}\text{J},\ 120.90\ (\overline{C}(4)_{\text{HetAr}}\text{J},\ 119.28\ (C(7)_{\text{HetAr}}\text{J}),\ 116.31\ (C_{Ar}-C\text{H}_3),\ 107.28\ (C_{Ar}-COO-),\ 68.99\ (Ar-C\text{H}_2O-),\ 61.02\ (Ar-OC\text{H}_3),\ 34.52\ (-C\text{H}_2-CON\text{H}-),\ 34.50\ (-C\text{H}_2-C\text{H}_2-),\ 22.89\ (Ar-C\text{H}_2-),\ 16.49\ (=C(-C\text{H}_3)-),\ 11.41\ (Ar-C\text{H}_3);\ R_{T,HPLC}=\ 13.238\ \text{min},\ \text{purity}\ 97.60\%;\ \text{HRMS}\ (m/z)\ calculated\ for\ C_{24}\text{H}_{23}\text{N}_3\text{O}_7\text{S}\ [M\ -\ H]^-:\ 496.1178;\ found:\ 496.1442.$





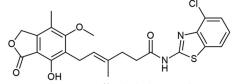
Method A - 1.0 eq DIPEA, washing step limited to cold ACN usage, afforded the product as white solid with 66% yield; mp. 165–168 °C; ¹H NMR (DMSO–d₆) δ 12.35 (s, 1H, –NH–), 9.34 (s, 1H, Ar–OH), 7.47 (d, J = 7.9 Hz, 1H, CH_{HetAr}), 7.24 (t, J = 8.0 Hz, 1H, C(6)H washing step limited to cold ACN usage HetAr), 6.99 (d, J = 8.0 Hz, 1H, CH_{HetAr}), 5.17 (t, J = 6.7 Hz, 1H, >C = CH–), 5.03 (s, 2H, Ar-CH₂O-), 3.91 (s, 3H, HetAr-OCH₃), 3.64 (s, 3H, Ar-OCH₃), 3.28 (d, J=6.9 Hz, 2H, Ar-CH₂-), 2.59 - 2.51 (m, 2H, -CH₂-), 2.29 (t, J = 7.4 Hz, 2H, -CH₂-), 1.95 (s, 3H, Ar-CH₃), 1.78 (s, 3H, -CH₃); ^{13}C NMR (DMSO–d_6) δ 171.92 (–CONH–), 170.67 (–COO–), 163.01 (C_{Ar}-OMe), 156.70 (C(2)_{HetAr}), 153.11 (C_{Ar}-OH), 152.35 (C(4)_{HetAr}), 145.98 (\underline{C}_{Ar} -CH₂O-), 138.94 (C(3a)_{HetAr}), 133.66 (>C = \overline{C} (-CH₃)-), 133.38 (C(7a)_{HetAr}), 124.73 (C(6)_{HetAr}), 123.71 (C_{Ar}-CH₂-CH=), 122.59 ($>\overline{C} = C(-CH_3)-$), 116.31 ($C_{Ar}-CH_3$), 113.85 ($C(\overline{7})_{HetAr}$), 108.31 (C(5)_{HetAr}), 107.27 (C_{Ar}-COO-), 69.00 (Ar-CH₂O-), 61.02 (Ar-OCH₃), 56.38 (HetAr–OCH₃), 34.70 (–CH₂–CONH–), 34.46 (–CH₂–CH₂–), 22.88 (Ar-CH₂-), 16.44 (=C($-CH_3$)-), 11.35 (Ar-CH₃); $R_{T,HPLC}$ = 12.274 min, purity 98.12%; HRMS (m/z) calculated for C₂₅H₂₆N₂O₆S [M — H]⁻: 481.1433; found: 481.1695.

N-(4-methylbenzo[d]thiazol-2-yl) mycophenolate (A17).



Method A – 1.0 eq DIPEA, washing step limited to cold ACN usage, afforded the product as white solid with 67% yield; mp. 173–176 °C; ¹H NMR (DMSO–d₆) δ 12.32 (s, 1H, –NH–), 9.31 (s, 1H, Ar–OH), 7.74 (d, J=7.7 Hz, 1H, CH_{HetAr}), 7.25 (d, J=7.2 Hz, 1H, CH_{HetAr}), 7.20 (t, J=7.5 Hz, 1H, C(6)H_{HetAr}), 5.18 (t, J=6.8 Hz, 1H, >C=CH–), 4.95 (s, 2H, Ar–CH₂O–), 3.65 (s, 3H, Ar–OCH₃), 3.27 (d, J=6.9 Hz, 2H, Ar–CH₂–), 2.60–2.52 (m, 5H, –CH₂– and HetAr–CH₃), 2.29 (t, J=7.3 Hz, 2H, –CH₂–), 1.93 (s, 3H, Ar–CH₃), 1.78 (s, 3H, –CH₃); ¹³C NMR (DMSO–d₆) δ 172.11 (–CONH–), 170.56 (–COO–), 168.41, 162.78 (C_{Ar}–OMe), 157.39 (C(2)_{HetAr}), 153.01 (C_{Ar}–OH), 147.95 (C(3a)_{HetAr}), 130.17 (C_{HetAr}), 126.93 (C_{HetAr}), 123.75 (C_{HetAr}), 123.67 (C_{Ar}–CH₂–CH=), 122.61 (>C=C(–CH₃)–), 119.38 (C(7)_{HetAr}), 116.16 (C_{Ar}–CH₃), 107.19 (C_{Ar}–COO–), 68.83

N-(4-chlorobenzo[d]thiazol-2-yl) mycophenolate (A18).



Method A - 1.0 eq DIPEA, afforded the product as chalky solid with 76% yield; mp. 223–227 °C; ¹H NMR (DMSO–d₆) δ 12.64 (s, 1H, -NH-), 9.33 (s, 1H, Ar-OH), 7.93 (d, J=7.9 Hz, 1H, CH_{HetAr}), 7.52 (d, J = 7.7 Hz, 1H, CH_{HetAr}), 7.30 (t, J = 7.9 Hz, 1H, C(6)H_{HetAr}), 5.18 (t, J=6.8 Hz, 1H, >C=CH-), 5.01 (s, 2H, Ar-CH₂O-), 3.64 (s, 3H, Ar-OCH₃), 3.28 (d, J=6.9 Hz, 2H, Ar-CH₂-), 2.62 - 2.53 (m, 2H, $-CH_2-$), 2.29 (t, J = 7.3 Hz, 2H, $-CH_2-$), 1.94 (s, 3H, Ar $-CH_3$), 1.78 (s, 3H, –CH₃); ¹³C NMR (DMSO–d₆) δ 172.48 (–CONH–), 170.53 (-COO-), 162.79 (C_{Ar}-OMe), 159.34 (C(2)_{HetAr}), 153.05 (C_{Ar}-OH), 145.99 (C_{Ar} -CH₂O-), 145.77 (C(3a)_{HetAr}), 133.48 (>C = C(-CH₃)-), (C_{Ar}-COO-), 68.85 $(Ar - CH_2O -), 60.95$ $(\overline{Ar} - OCH_3),$ 34.68 $(-CH_2-CONH-)$, 34.27 $(-CH_2-CH_2-)$, 22.87 $(Ar-CH_2-)$, 16.44 $(=C(-CH_3)-)$, 11.33 (Ar-CH₃); $R_{T,HPLC} = 13.969$ min, purity 97.67%; HRMS (m/z) calculated for C₂₄H₂₃ClN₂O₅S [M - H]⁻: 485.0938; found: 485.1194.

Biological evaluation

Research materials

T-Jurkat (Clone E6-1, ATCC TIB-152), a human T lymphoblastoid cell line from an acute T-cell leukaemia was received from a cell bank of Department of Medical Immunology, Medical University of Gdansk. Prior to testing, cells were thawed and passaged for approximately 7 days in order to enter the logarithmic growth phase.

Human peripheral blood mononuclear cells (PBMCs) were obtained from buffy coats of anonymous, healthy volunteers, with the average age 33 ± 6 years, from the Regional Centre for Blood Donation and Treatment in Gdansk.

Growing medium

T-Jurkat cell line was suspended in a complete RPMI medium containing RPMI-1640 media (Sigma Aldrich, Poznań, Poland), 10% FBS (Sigma Aldrich, Poznań, Poland), penicillin (p, 100 U/mL) and streptomycin (s, 100 mg/mL, Sigma Aldrich, Poznań, Poland). PBMCs were suspended in a complete X-VIVO medium containing X-VIVO20 media (X-VIVO 20, Lonza, Verviers, Belgium), 10% heatinactivated human AB serum (National Blood Bank, Raciborz, Poland), penicillin (p, 100 U/mL) and streptomycin (s, 100 mg/mL).

MPA derivatives preparation

To prepare appropriate concentrations of tested compounds, the substances were dissolved in DMSO (Sigma Aldrich, Poland) with 1% addition of Tween20 to obtain final concentration of 0.01 M. The solution was diluted 10-fold and then 5-fold in appropriate medium without additives (RPMI-1640 for T-Jurkat cell line; X-

VIVO20 for PBMCs) to obtain the following concentrations: 200; 20; 2; 0.2; 0.02 μ M.

PBMCS isolation

PBMCs were isolated by gradient centrifugation, as described previously⁵². Buffy coats were diluted with phosphate-buffered saline (PBS, pH 7.4, ThermoFisher Scientific, Waltham, MA) in a 1:1 ratio, layered at Ficoll-Paque[™] PLUS (VWR International, Gdańsk, Poland) and centrifuged. PBMCs layer was collected and washed twice with PBS. The cells were counted and suspended in the appropriate medium depending on their purpose.

In vitro cytotoxicity assay with XTT dye

The cytotoxic effect of derivatives of MPA was measured using in vitro 2,3-bis(2-methoxy-4-nitro-5-sulfophenyl)-2H-tetrazolium-5carboxanilide (XTT)-based assay (Roche, Basel, Swiss). PBMCs suspended in complete X-VIVO20 medium and T-Jurkat cells suspended in complete RPMI-1640 medium were added on a 96-well plate at a concentration of 5×10^5 cells/50 µL/well. 50 µL of examined compounds, prepared as described in MPA derivatives preparation section) were added to experimental wells to obtain final concentrations: 100; 10; 1; 0.1; 0.01 µM. As a background control we used medium without cells, as a positive control we used cells untreated with compounds. After 48 h for T-Jurkat cells and 72 h for PBMCs of incubation in a humidified atmosphere with 5% CO₂ at 37 °C, 50 µL of XTT reagent (freshly prepared according to the manufacturer's procedure) was added to all samples. Plates were incubated for 24 h (5% CO2 at 37 °C). The conversion of water-soluble yellow tetrazolium XTT salt into orange formazan was monitored by measuring the optical density (OD) at 450 nm on Agilent BioTek Epoch Microplate Spectrophotometer (Agilent Technologies, Santa Clara, CA), with a background wavelength at 690 nm. Data were expressed as IC_{50} to compounds that inhibited more than 50% of the cell viability. Each sample was performed in triplicates. The final concentration of DMSO in the culture did not influence cellular viability. Cell viability was determined at the highest concentration of the tested compound. For this purpose, TC20 cell counter (Biorad, Hercules, CA) was used with trypan blue exclusion.

In vitro dye-based proliferation assay

PBMCs and T-Jurkat cells were washed twice with warm PBS (37 °C), suspended in fresh PBS at a final concentration of $10-30 \times 10^6$ cells/mL and labelled with $1\,\mu$ L of VPD450 (BD Biosciences, Franklin Lakes, NJ) at 37 °C for 15 min. Each 5 min cells were vortexed to provide uniform staining^{42,43}. Subsequently, cells were washed with PBS and then with complete medium (X-VIVO20 for PBMCs and RPMI-1640 for T-Jurkat cells). After this step, cells were suspended in fresh complete medium, seeded on 96-well plates (1×10^6 cells/well). PBMCs were additionally stimulated with magnetic beads coated with anti-CD3 and anti-CD28 antibodies (Invitrogen, Carlsbad, CA) in 1:0.5 cell:bead ratio. Of 100 µL of examined derivatives of MPA were added to experimental wells to receive final concentrations: 100; 10; 1; 0.1; 0.01 µM. In parallel, stained and not stimulated cells, as well as stained and stimulated cells, were seeded in the same concentration as controls. After 48 h for T-Jurkat and 72 h for PBMCs, cells were analysed with flow cytometer LSRFortessa (BD Biosciences, Franklin Lakes, NJ) and Kaluza C Flow Cytometry Analysis Software (Beckman Coulter, Brea, CA). Data were expressed as EC₅₀ to compounds that inhibited more than 50% of the cell proliferation. Cell

viability was determined at the highest concentration of the tested compound. For this purpose, TC20 cell counter (Biorad) was used with trypan blue exclusion.

Flow cytometry Annexin V/PI assay

The percentage of apoptotic and necrotic cells after treatments were detected using an FITC Annexin V Apoptosis Detection Kit I (BD Biosciences). According to the manufacturer's protocol, cells were washed with cold PBS and then resuspended with 1XBinding Buffer at a final concentration of 1×10^6 cells/mL. To 1×10^5 of cells 5 µL of Annexin V-FITC and 5 µL of propidium iodide (PI) were added. After 15 min at room temperature in the dark, flow cytometry analysis was performed. We divided cells into four groups according to Annexin V/PI results: living cells (Annexin V-negative, PI-negative); early apoptotic cells (Annexin V-positive, PI-positive, PI-positive, PI-positive, PI-positive) and necrotic cells (Annexin V-negative, PI-positive). Cells were analysed with flow cytometer LSRFortessa and Kaluza C Flow Cytometry Analysis Software.

Statistical analysis

Statistical analysis was performed using GraphPad Prism version 9 (GraphPad Software Inc., La Jolla, CA). Two-tailed unpaired Student's *t*-test was used to evaluate differences between MPA and new derivatives of MPA; differences were considered as statistical significant when p < 0.05. EC₅₀ and IC₅₀ values were calculated by 4-parametric non-linear regression using GraphPad Prism version 9. Significance was calculated using the *t*-test, significant results are marked with *(p < 0.05), **(p < 0.01), or ***(p < 0.001).

Acknowledgements

NMR studies were performed in the Nuclear Magnetic Resonance Laboratory of Gdańsk University of Technology.

Disclosure statement

The authors do not report any competing financial interests or personal interrelations which could have governed publication's outcomes.

Funding

Financial support for this work was supported by the internal grant DS035136 of Gdańsk University of Technology and statutory grant no. 02–10022/272 of Medical University of Gdańsk.

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